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Original Research Article

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5 Abstract

Obesity is associated with an increased risk of metabolic syndrome such as type 2 diabetes, 6 7 insulin resistance, dyslipidaemia and non-alcoholic fatty liver disease. Obesity occurs as a result of imbalance between food intake and energy expenditure leading to excessive 8 accumulation of adipose tissue. NAFLD is the most common liver condition and related to 9 resistance of insulin. Insulin resistance is associated with increase influx of lipid into the liver 10 promoting accumulation of hepatic triglyceride. The aim of this study is to develop an 11 12 experimental model of hepatic steatosis with lipid over accumulation. HepG2 cells were cultured for 24 hours in free fatty acid media (1:2 palmitic acid and oleic acid respectively). 13 14 Intracellular lipid content and lipotoxicity were determined by oil red O staining followed by colorimetric detection. This experiment was accomplished by defining the experimental 15 conditions of lipid exposure that leads to significant intracellular fat accumulation in the 16 17 absence of lipotoxicity with 1 mM of free fatty acid media. As a result, oleic and palmitic acids could be over accumulated in HepG2 cells. 1 mM free fatty acid media did not affect 18 the cell integrity and did not cause lipotoxicity of the cells. 19 20

AN EXPERIMENTAL MODEL OF HEPATIC STEATOSIS

TO DETECT LIPID ACCUMULATION

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Key words: hepatocellular carcinoma (HepG2) cells, fat accumulation, Oil Red O,
 Nile Red, lipotoxicity, ...

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25 1. INTRODUCTION

26 Increased incidence in obesity is reaching epidemic proportions because of lifestyle modification especially change in eating habits. Obesity complications including type 2 27 28 diabetes, cardiovascular diseases, metabolic disorders and many types of cancer [1] have raised a significant health concern in the world especially in developed countries [2,3]. 29 Therefore, many studies of molecules regulating the development of obesity and its 30 associated pathologies is ongoing to treat and prevent obesity [4]. An important link between 31 obesity, hypertension and sympathetic nerve activity (SNA) is leptin [5]. Leptin is a 16 KDa 32 protein produced by adipocytes and controls body weight by regulating appetite and energy 33 expenditure [6,7]. Absence of functional leptin receptors or deficiency in leptin is related 34 35 with distinct hyperphagia and reduced energy expenditure [8], whereas viral vectors increase 36 leptin by overexpression of leptin gene or infusion of leptin, this lead to decrease food intake 37 and increases expenditure of energy [9,10]. Leptin (adipocyte hormone) increases with increase proportion to adipose tissue mass and decrease with weight loss [11] and plays a key 38

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role in glucose homeostasis facilitated by its direct action on the central nervous system [12]. 39 40 Glucose is an important nutrient and source of energy whose homeostasis is important to maintain proper cell functions, since the physiology of the body can be weakened by either 41 42 hypoglycaemia or hyperglycaemia leading to cell death. The inability of the cells to use or 43 take up glucose as an energy source upon stimulation by insulin is defined as insulin resistance (IR) [13]. The intrinsic protein tyrosine kinase activity is induced by binding of 44 45 insulin to its receptor; the tyrosine residues are phosphorylated in the insulin receptor as a result of activating intrinsic protein tyrosine kinase activity this is the first step. "Subsequent 46 steps involved the activation of the heterodimeric p85/p110-PI3K complex including 47 generation of the lipid second messenger PIP3 (Phosphatidylinositol (3, 4, 5)-trisphosphate) 48 which activates PDPK1 (phosphoinositide-dependent protein kinase-1), PKB/Akt (Protein 49 Kinase B) and a typical PKC isoform and recruitment of adapter molecule's insulin receptor 50 substrates (IRS), IRS1 and IRS2" [14,15]. 51 The most common form of chronic liver disease is non-alcoholic fatty liver disease 52

(NAFLD), its prevalence increase with increase incidence of obesity [1,16]. NAFLD is 53 defined as excess accumulation of fat in the liver that is not a result of alcohol consumption, 54 genetic disorders or drug use [17]. The accumulation of lipids in micro and macro vesicles in 55 56 more than 5 % hepatocytes, mostly in the perivenular hepatocytes is known as Non-alcoholic steatohepatitis (NASH). NASH associated with steatosis and necro inflamation is a more 57 severe form of NAFLD and may lead to hepatic fibrosis and cirrhosis [18]. Progession of 58 simple steatosis to NASH is contributed by increased delivery of fatty acids to the liver. Fatty 59 acids such as palmitic and oleic acid are found in triacyl-sn-glycerols of seed oils and animal 60 depot fats in our daily diet. Palmitic acid is a saturated fatty acid that induces apoptosis in 61 62 hepatocytes [19]. In the presence of oleic acid, the palmitate-induced apoptosis is reduced [19]. It is known that lipid accumulation in NASH is induced by free fatty acids (FFA) [20]. 63 To study NAFLD, HepG2 cells have been widely used. HepG2 cells are derived from tumor 64 cells, thus they behave differently from normal cells [21]. Lipotoxicity - a metabolic 65 syndrome caused by the accumulation of free fatty acids in non-adipose tissue can lead to 66 cellular dysfunction and eventually inevitable death [22]. Triglycerides is the major lipid 67 stored in NAFLD recent data suggest that triglyceride accumulation may be hepatoprotective 68 in the liver. A diacylglycerolacyltransferase 1 or 2 catalyze the synthesis of hepatic 69 70 triglyceride [23]. Accumulation of FFA is likely to have toxic effects in hepatocytes in 71 contrast to triglyceride. Li et al. (2009) demonstrates the main determinant of hepatocellular damaged in NAFLD is the ratio of monounsaturated FAs (MUFA) to saturated FAs (SFA) 72 73 [24].

Protein tyrosine phosphatases (PTPs) have been involved in the modulation of glucose homeostasis *in vivo*, including protein tyrosine phosphatase 1B (PTP1B) [25]. The IR PTK in liver and muscle is dephosphorylated by PTP1B to regulate glucose homeostasis. Increasing the expression of PTP1B leads to insulin resistance in people and rodents and knockout of PTP1B is associated with leanness and insulin sensitivity in rodents, this suggests that PTP1B is an important molecular target for the treatment of diabetes and obesity. Due to the role of Comment [T16]: Binding of insulin induces the

intrinsic protein tyrosine kinase activity to its

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80 PTP1B in IR and after leptin signalling, PTP1B can be used as a target for the development

of therapeutics for treatment of obesity and type 2 diabetes. Antisense oligonucleotides that

target PTP1B are in clinical trials whereas drugs inhibiting the activity of PTP1B are in

preclinical development [26]. Decreasing the level of PTP1B in peripheral tissues is

84 associated with improved insulin sensitivity, regulates hyperglycaemia and reduced obesity in

these mice [27]. This suggests that PTP1B plays an important role in regulation of body mass

in one or more peripheral tissues independent on effects in the brain [27]. A common model

to study hepatic steatosis is FFA induced lipid accumulation in hepatocytes [28]. The aim of

this study is to develop an experimental model of hepatic steatosis using HepG2 cell line and

test the lipid accumulation in liver cells in the designed model. To obtain the aim, the study
 used Oil red O and Nile red staining to measure lipid levels in HepG2 cell lines.

91 2. MATERIALS AND METHODS

92 2.1. Materials

Human hepatocellular carcinoma (HepG2) cells were supplied by Sheffield Hallam
University. Cells were grown in high glucose (4.5 g/L) Dulbecco's modified Eagle's medium

95 with ultra-glutamine (DMEM) (Lonza, UK), 10 % Fecal Calf Serum (FCS) (Gibco, UK) and

96 1 % penicillin/streptomycin (Lonza, UK).

Basic culture media was prepared with 200 ml of DMEM (Lonza, UK), 2 ml 1 %
penicillin/streptomycin (Lonza, UK) and 2 g of 1 % Bovine Serum Albumin (BSA) (ThermoFisher).

Oleic acid (Sigma-Aldrich) and palmitic acid (Sigma-Aldrich) dissolved in dimethyl
sulfoxide were diluted in basic media to obtain 30 mM of free fatty acid (FFA) media (stock
solution). This stock solution was further diluted with basic culture media to obtain 1 mM
FFA media.

104 2.2. Oil red O colorimetric assay

105 Stock solution of Oil red O (ORO) was prepared by dissolving 0.7 g of ORO in 200 ml of

106 Isopropyl alcohol. Bakers' formalin was prepared by adding 10 ml of 37 % formaldehyde and

107 10 ml of 10 % (w/v) calcium chloride solution to 80 ml of water and stored at 4°C. To prepare

108 glycerol gelatine, 5 g gelatine was gently mixed with 50 ml glycerol in 50 ml water at 50°C,

and stored at 4°C, then heating at 55-60°C before use.

110 Cells were seeded in 12-well plate for 24 hours before treatment with 1 mM fatty acid at a

density of 50000 cells per well. A half of the plate (6 wells) was used as control samples and

the other half was treated with 1 mM of FFA and incubated for 24 hours at 37°C and 5 %

113 CO₂, in SANYO incubator before staining with Oil red O.

114 A 60 % ORO solution was prepared by diluting stock ORO with water at 3:2 ratio. Harris

haematoxylin (Sigma Aldrich, UK) and 60 % ORO were filtered before use. Treatment and

basic media were removed from each well and cells were washed 2 times in Hanks Balanced

117 Salt solution (Sigma Aldrich, UK), the excess was poured off and blotted with a dry tissue.

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118 Each well was fixed with 780 μ l of bakers' formalin at 4°C for 30 minutes, the excess was

poured off and blotted with a dry tissue. 780 μl of 60 % ORO was added to each well and left

to incubate for 10 minutes at room temperature. Cells were rinsed with water for 5 minutes

after staining. Excessive water was blotted with dry tissue before staining cells with 780 μ l of

Harris haematoxylin and incubated for 2 minutes at room temperature. Cells were rinsed with
water for 5 minutes. Excessive water was blotted with dry tissue and 2 drops of glycerol

124 gelatine were added to each well. Stained cells were observed under x400 magnification of

the microscope and quantified using Image J program. Threshold was set at 163 and 213 for

126 minimum and maximum, respectively.

127 2.3. Proliferation Assay

128 100x stock solution of alamar blue (Sigma-Aldrich, UK) was prepared by dissolving 1 mg

129 alamar blue in 1 ml of 1x PBS. Working concentration was achieved by diluting 30 μ l of

130 100x stock solution in 30 ml of basic media. To measure cell proliferation, negative control

131 was achieved by adding 1 ml of 1 % Triton X-100 solution to each control and treated

132 sample, then incubated for 1 hours before staining with Alamar Blue. After incubation of the

negative control, all cell media was removed and 2 ml of alamar blue solution was added

followed by a 4-hour and 20-hour incubations at 37°C and 5 % CO₂, in SANYO incubator.

Absorbance was measured at 570 nm using GENESYS 10S UV-VIS (Thermo-Fisher) after 4and 20 incubation hours.

137 2.4. Protein quantification

138 Cells were seeded in a 6-well plate for 24 hours before treatment with 1 mM fatty acid at a seeding density of 50000 cells per well. Cells were incubated at 37 °C and 5 % CO2, in 139 SANYO incubator. Half (3) of the plate was used as control wells and the other half was 140 treated with 1 mM FFA for 24 hr at 37°C and 5 % CO₂, in SANYO incubator. Bradford 141 reagent was prepared by dissolving 50 mg Coomassie brilliant blue in 24 ml 99 % ethanol; 50 142 ml of 85 % (w/v) phosphoric acid was added and diluted to a final volume of 500 ml with 143 water, this solution was filtered before use. Protein standard was made by dissolving 0.05 g 144 BSA in 50 ml warm water at a concentration of 1000 µg/ml. A serial dilution was prepared in 145 the following concentrations of 1000, 500, 250, 125, 63, 31 and $0 \mu g/ml$. 146

After treatment, basic media was removed from each well and cells were washed with 1x 147 PBS. 20 µl of cell lytic M reagent was added to each well and incubated at room temperature 148 for 15 minutes on a shaker. Lysed cells were collected by scraping. The lysed cells were 149 centrifuged for 15 minutes at 12000 x g at room temperature. The supernatants were removed 150 and the pellets were re-suspended in 100 μ l of water. Each pellet and supernatant sample 151 152 were diluted 1:10 with water, 10 µl was pipetted in triplicate into a flat bottomed 96-well 153 plate. Standards were also added to the plate in duplicate. 200 µl of Bradford reagent was 154 added into each well and absorbance was measured by MULTISCAN FC; Thermo-Fisher, 155 UK at 595 nm.

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156 2.5. Nile Red fluorescence assay

157 Cells were seeded in a 6-well plate for 24 hours before treatment with 1 mM fatty acid at a 158 seeding density of 50000 cells per well. Cells were incubated at 37°C and 5 % CO₂, in 159 SANYO incubator. The half (3 wells) of each plate was used as control wells and the other 160 half was treated with 1 mM concentration for 24 hours at 37°C and 5 % CO₂, in SANYO 161 incubator. Working solution of Nile Red (1 μ g/ml) was achieved by diluting 1 mg in 1 ml 162 methanol before diluting 5 μ l with 5 ml 1x PBS.

163 Cells were washed with 500 µl of PBS (Lonza, UK). 250 µL of trypsin was added to each 164 well and placed in an incubator for 5 min for the cells to dislodge. 250 µl of media was added 165 to each well and pipetted into different tubes. Cells were centrifuged at 500 x g for 5 minutes. 166 Supernatants were discarded and pellets were re-suspended in 1.5 ml 1xPBS and centrifuged at 500 x g for 5 minutes. 500 µl of 1x Nile Red in PBS was added and incubated for 5 167 168 minutes on ice. Cells were washed in 1.5 ml PBS, supernatant was discarded and pellet was 169 re-suspended in 500 µl of PBS. Nile red fluorescence was determined by flow cytometry with Becton Dickinson FACS Calibur System, FL2 emission channel, at excitation wavelength of 170 171 488 nm and emission wavelength of 550 nm.

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173 **3. RESULTS**

174 3.1. Oil red O colorimetric assay

After 24-hour incubation with 1 mM of FFA media in a 12-well plate, cells were stained with Oil red O to observe the accumulation of lipid (palmitic acid and oleic acid) at x400

magnification of the microscope. Compared with the control samples, the appearance of red

spots in Figure 1B shows lipid accumulation in FFA-treated HepG2 cells. The quantification

179 of stained HepG2 cells using Image J is expressed as % average areas of 6 wells. In control

cells, the percentage area was approximately 12% while that was nearly 16% in treated cells.



Figure 1. Oil Red O staining of HepG2 cells in A) control samples, B) FFA-treated HepG2 cells samples * red arrow shows lipid droplet; black arrow shows cell nucleus

185 3.2 Proliferation Assay

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186 Proliferation assay was used to monitor the response and health of HepG2 cells in culture

after treatment with 1 mM FFA. The assay shows the lipotoxicity of HepG2 cells. Treatment

188 of HepG2 cells with 1 mM of FFA media did not significantly decrease the cell viability,

189 compared to control cells. At the beginning point, the reduction of alamar blue calculated was

190 low, under 5% in both control and treated cells. After incubation, this number increased

191 following the increase in incubation time. After incubation in 4 hours, the amount of alamar

blue that was reduced in both samples was 17 %. This proportion reached 52 % and 57 % for

193 control and treated cells, respectively after 20-hour incubation. The cell growth inhibition

194 was difference in treated and control HepG2 cells. After incubating these cells in 4 hours, the

difference percentage was 95 % and increased up to 112 % after 20-hour incubation.

The amount of protein has been successfully measured in HepG2 cells. In the standard curve, the absorbance increased when increasing the concentration of protein from 0 to 31, 63, 125,

198 250, 500 and 1000 μ g/ml. The regression equation was A = 0.0012C + 0.2698 with the (R² = 0.9962). Table 1 shows the difference in concentration of proteins extracted from pellet and suspension of the samples, and difference between untreated and treated cells. High protein

concentrations were seen in the suspension of both the control and treated cells. The pellet ofcells had low protein concentration.

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Table 1. Concentration of protein calculated from the standard curve

Samples		Protein concentration (µg/ml) in triplicate		
Control colla	pellet	0.54	0.51	0.45
Control cens	suspension	2.01	2.09	1.83
T	pellet	0.62	0.81	0.59
I reated cens	suspension	1.69	1.87	1.09

204 3.3. Nile red fluorescence assay

205	The content of intracellular lipid droplets was determined by Nile red staining and the cellular
206	FFA uptake was quantified by flow cytometric assay. Cells exposed to 1 mM FFA for 24
207	hours induced fat accumulation. Unstained control cells (figure 2A) were used to adjust the
208	settings of the flow cytometer to measure intensity of unstained treated cells. This procedure
209	is similar to measure intensity of stained treated cells. Unstained cells gave very low signals
210	as compared with stained cells. For example, unstained treated cells gave only 2.75 while
211	stained treated cell gave up to 6130 intensity unit. Compared to stained control cells (figure
212	2G) which caused only 1366 intensity, the fluorescence signal of lipid in the stained treated
213	cells was evidently higher, up to roundly 6130 intensity unit (figure 2H). In summary, the
214	difference in fluorescence intensity shows an increase in intracellular lipid accumulation in
215	cells treated with FFA media.

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223 DISCUSSION

NAFLD causes chronic liver disease and prevalent increase with growing epidemic of obesity
worldwide. The prevalence and impact of NAFLD on the growing epidemic of obesity make
NAFLD have become the most common cause of liver disease [29]. NAFLD is associated
with several non-hepatic related complications and has the potential to develop to hepatic
fibrosis and end-stage liver disease [30]. There is a general understanding that patients with
NASH have increased lipolysis and subsequently a high circulating FFA level, accumulation

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of lipid in liver cells is contributed by increase in FFA inflow and de novo lipid synthesis

- [31]. A model for investigating NASH that is usually used is incubating hepatocytes with
- FFA [28]. The structure of fatty acids affects their activity biologically and they are classified
- 233 as saturated and unsaturated fatty acids. The saturated FAs induce lipotoxicity, insulin
- resistance and apoptosis. A saturated fatty acid that induces apoptosis in liver cells is palmitic
- acid. Oleic acid is discovered to be more steatogenic but less apoptotic than palmitic acid
 [19,32]. Co-incubation of liver cells with both palmitic and oleic acid results in higher
 amount of fat accumulation than incubation with only palmitic acid. This co-incubation also
- 238 lowers palmitate-mediated apoptosis, indicating protective feature of oleic acid [19,32]. In
- this study, HepG2 cells were incubated with a mixture of PA and OA (1:2) respectively, to
- 240 develop a model to test the effect of PTP1B inhibitor on gluconeogenesis in vitro. To
- 241 determine if HepG2 cells accumulate lipid, Oil red O was performed, flow cytometry based
- on Nile red staining was performed to determine the intracellular lipid content.

243 With this experiment, the quantification with ImageJ shows that HepG2 cell line can 244 accumulate FFA, which was stained by ORO and viewed under microscope x 400 magnification. The cells were cultured in high glucose media, some lipids were seen in 245 control but fewer in number compared to the treated cells. This result is similar with the study 246 247 of Chavez-Tapia et al. (2011) [33], reaching the conclusion that hepatic cell line (HuH7) and no tumoral immortalized human hepatocytes (IHH) used could accumulate FFA, and the 248 increase in fat content was not related to critical deterioration of the cell integrity, in 249 250 accordance with clinical and in vivo experimental information.

251 Yao, et al., 2011 demonstrate that 1 mM concentration of FFAs caused fat accumulation but not lipotoxicity [34]; however, group of HepG2 cells treated with 2 and 3 mM FFAs 252 253 significantly increased the lipotoxicity in cells. The proliferation assay was used to determine 254 how proliferative HepG2 cells are and to confirm if the concentration of FFA causes lipotoxicity in this study. In this study, after incubation in 4 hours, there was 17% reduction 255 of alamar blue in both the control and treated cells, while 20-hour incubation gave 52% and 256 257 57% reduction of alamar blue in control and treated wells, respectively. This means that the cells are dividing rapidly in 20 hours than 4 hours because of long period of incubation. 1 258 259 mM FFA concentration showed no significant difference in the proliferation of the treated 260 HepG2 cells compared to the control cells; this indicates that 1 mM FFAs cause's lipid accumulation but not cell toxicity. This is similar to previous study [34] on lipotoxicity in 261 262 HepG2 cells that support the findings in this practical. In 4-hour incubation, treated cells are differentiating at 95% of the rate of control cells, this indicate that the treated cells have 263 reduced alamar blue by 95% of what the control had done or growth in treated is inhibited by 264 265 5% compared to control. However, in 20-hour incubation, treated cells are differentiating at 112% of the rate of control cells, this indicates that there is 0% growth inhibition in the 266 treated cells compared to control cells. 267

To determine the intracellular lipid droplets contents, HepG2 cells were stained with Nile red.
Exposure of cells to 1 mM FFA for 24 hours induced accumulation of lipid. Figure 4 (A and

B) shows the forward and side scatter of the unstained control and unstained treated cells

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respectively. The forward scatter is used to identify the size of cells; it measures the size and 271 272 shape of the cells. The side scatter is used to determine the complexity of the internal environment of the cell. Cells with a large amount of complexity in the internal environment 273 274 will have a higher side scatter signal than cells with no or low complexity [35]. In figure 4 (A and B), the forward scatter shows a single cell population that are exactly the same size 275 (HepG2 cells). Figure 4B has a higher side scatter than figure 4A, which shows figure 4B 276 277 cells have more internal complexity than cells in figure 4A as a result of treating cells with 1 mM FFAs. In figure 4C and 4D, the peak was on the left side of the axis (no fluorescence) 278 indicating the cells were not stain with Nile red. Microscopic image shows the presence of 279 cytoplasmic lipid droplets. This data was confirmed by flow cytometry in figure 5. The 280 highest peak was observed in figure 5D (treated stained) compared to figure 5C (control 281 stained) due to fat accumulation in the cells. When the cells were treated with 1 mM FFA the 282 peak shifts to the right, this corresponds to the significant increase in the geometric mean 283 284 fluorescence intensity which is 6x more compared to the control stained cells. This is similar 285 to previous study [34] that shows an increase in fluorescence intensity of lipid in treated 286 stained cells compared to control stained cells under the microscope, they further showed the 287 difference of fluorescence intensity in treated cells compared to control cells by flow cytometry, they observed a dose dependent increase in lipid accumulation in cells treated 288 with FFA. Nile red attached to the lipids in the treated cells could make a more complex 289 290 environment in the cells because of the lipid droplets. The more complex the internal environment of the cell was, the higher the side scatter signal was. This is why the fluorescent 291 292 intensity of the treated cells was higher 5 times compared to the control stained cells. 293 Bradford assay was used to measure the concentration of protein in a solution, which was proposed by the study of Cheng et al. (2016) [36]. To normalize the concentration of protein 294 295 for western blotting, Bradford assay was performed. As a result, the highest protein 296 concentration was seen in control suspension while the lowest concentration of protein was in 297 control pellet.

298 CONCLUSION

299 PTP1B is an important molecular target for the treatment of diabetes and obesity. Targeting this inhibitor to reduce the complications that are associated with obesity such as hepatic 300 301 steatosis and type 2 diabetes is of importance to the health care system. Some of the objectives of these study which include, identifying markers that are involved in 302 gluconeogenesis that may affect PTP1B, measuring protein concentration using western 303 304 blotting were not achieved due to time constraint. PTP1B inhibitor is supposed to be tested on the cell model that was designed but due to time constraint the inhibitor was not tested. If the 305 306 inhibitor was tested on the model in all the assays performed above (Oil red O, Nile red, proliferation assay and flow cytometry), it is assumed that the cells may not have accumulate 307 the FFA, lipotoxicity will not be seen and this will prevent non-alcoholic fatty liver disease 308 and complications associated with it. In conclusion, the results show that HepG2 cells are 309 induced to lipid over accumulation by mixtures of oleate and palmitate acids. 1 mM FFA did 310 311 not affect the cell integrity and did not cause lipotoxicity of the cells. FFAs should be used

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- 312 with different concentrations of FFAs to develop further the model. Methods such as MTT
- assay can be used to assess the FFAs cytotoxicity to HepG2 cells. Cells treated with FFAs
- and stained with Annexin V/propidium iodide should be assayed for apoptosis using flow
- 315 cytometry in further experiments.

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