## Supplementary Materials: Antidiabetic, Lipid Normalizing, and Nephroprotective Actions of the Strawberry: A Potent Supplementary Fruit

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Table S1. Animals	' food and	water intake	during i	nterventions.
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Groups	Feed Intake (gm)	Water Intake (mL)
Healthy control	22.3 ± 1.24 *	50.24 ± 1.93 **
NIC-STZ control	$44.1 \pm 0.52$	$138.6 \pm 1.64$
MET-treated group	$42.4 \pm 9.51$	94.19 ± 0.74 **
SWE	37.9 ± 9.56 *	87.13 ± 0.63 ***
SHAE	26.7 ± 0.28 **	80 ± 3.24 **
SAE	59.8 ± 1.144 **	70 ± 0.45 **

Results are represented as mean  $\pm$  SE. gm/rat/day (n = 6); \*  $p \le 0.05$ ; \*\*\*  $p \le 0.01$ ; \*\*\*  $p \le 0.001$  as compared to the negative control group (Dunnett Multiple Comparisons Test).

**Table S2.**The efficiency of qRT-PCR for relative quantification of mRNA. Efficiency is calculated from the slope of the curve as  $E = 10(-1/\text{slope})^{-1}$ .

Sr. No.	Target Genes	Symbol	Efficiency			
Fatty Acid Metabolism Genes						
1	Carnitine Palmitoyltransferase 1A	CPT 1A	121.22			
2	Malonyl CoA: ACP Acyltransferase	MCAT	110.17			
3	Acetyl-CoA Carboxylase Alpha	ACACA	110.17			
4	acyl-CoA Synthetase Long-chain family member 1	ACSL 1	110.17			
5	Fatty Acid Synthase	FASN	93.06			
6	Fatty Acid Binding Protein	FABP	101.30			
Transcription Factors						
1	Peroxisome Proliferator-Activated Receptor Gamma	PPAR-γ	100.92			
2	Sterol Regulatory Element-Binding Proteins	SREBP	104.75			
3	Nuclear Factor-κβ	NF-κβ	108.43			
Inflammatory Markers						
1	Tumor Necrosis Factors- $\alpha$	TNF-α	98.71			
2	Interluekin 6	IL6	96.84			

**Table S3.** Experimental design employed in the study.

Days	НС	NIC-STZ	MET	SWE	SHAE	SAE
1st day	I. p. injection of saline solution and citrate buffer	I. p injection of nicotinamide (110 mg/kg) and STZ (65 mg/kg)	I. p injection of nicotinamide (110 mg/kg) and STZ (65 mg/kg)	I. p injection of nicotinamide (110 mg/kg) and STZ (65 mg/kg)	I. p injection of nicotinamide (110 mg/kg) and STZ (65 mg/kg)	I. p injection of nicotinamide (110 mg/kg) and STZ (65 mg/kg)
22nd day	-	Confirmation of stable hyperglycemia	Confirmation of stable hyperglycemia	Confirmation of stable hyperglycemia	Confirmation of stable hyperglycemia	Confirmation of stable hyperglycemia
23rd day	-	-	Metformin (200 mg/kg body weight, Per os)	Strawberry water extract (2 g/kg body weight, Per os)	Strawberry hydro-alcoholic extract (2 g/kg body weight, Per os)	Strawberry alcohol extract (2 g/kg body weight, Per os)
30th, 36th, 43rd day		Intermittent blood collection				
50th day	Blood samples were collected, animals were sacrificed. Liver, pancreas, brain, and kidney were harvested					

HC: Healthy control, NIC-STZ: NIC-STZ treated control, MET: Metformin-treated control, SWE: Strawberry water extract, SHAE: Strawberry hydro-alcoholic extract, SAE: Strawberry alcohol extract.

**Table S4.** Primer sequences of selected RT-PCR genes.

Sr. No.	Target Genes	Primers	Sequence			
Housekeeping Gene						
1	Channel debards 2 mb combate debardus comass	Forward	AGTTCAACGGCACAGTCAAG			
	Glyceraldehyde-3-phosphate dehydrogenase	Reverse	TACTCAGCACCAGCATCACC			
Fatty Acid Metabolism Genes						
1	Comition Delivited to a second 1 A	Forward	CACTGATGAAGGAAGAAGAC			
	Carnitine Palmitoyltransferase 1A	Reverse	CCAGTCACTCACGTAATTTG			
	Malamal Co A. ACD A military of our co	Forward	AAAACTCTAGGCTCAATCAAC			
2	Malonyl CoA: ACP Acyltransferase	Reverse	GGATGTGTGTATTTATGCCC			
2	A cotal Co A Composizione Alpha	Forward	AGCAGTATTTGAACACATGG			
3	Acetyl-CoA Carboxylase Alpha	Reverse	CAGTTCCAAGAAGTAGAAGC			
4 acyl	acyl-CoA Synthetase Long-chain family member 1	Forward	ACATTATGAACGATTGCTCC			
	acyr-coa synthetase Long-chain family member 1	Reverse	GCATTACACACTCTACAACG			
5	Fatter Asid Cropthago	Forward	AAAAGGAAAGTAGAGTGTGC			
	Fatty Acid Synthase	Reverse	GACACATTCTGTTCACTACAG			
6	Fatty Acid Rinding Protein	Forward	TGGAGGGTGACAATAAAATG			
	Fatty Acid Binding Protein	Reverse	TCATGGTATTGGTGATTGTG			
	Transcription Fact	tors				
1 Perox	Paravisama Praliforator Activated Pacantar Camma	Forward	AAGACAACAGACAAATCACC			
1	1 Peroxisome Proliferator-Activated Receptor Gamma		CAGGGATATTTTTGGCATACTC			
2 Sterol Reg	Sterol Regulatory Element-Binding Proteins	Forward	AAACCTGAAGTGGTAGAAAC			
	Sterof Regulatory Element-Dinality Proteins	Reverse	TTATCCTCAAAGGCTGGG			
3	Nuclear Factor 168	Forward	AAAAACGAGCCTAGAGATTG			
	Nuclear Factor-κβ	Reverse	ACATCCTCTTCCTTGTCTTC			
Inflammatory Markers						
1	Tumor Necrosis Factors- $\alpha$	Forward	CTCACACTCAGATCATCTTC			
1	Tumor recrusis ractors-a	Reverse	GAGAACCTGGGAGTAGATAAG			
2	Interleukin 6	Forward	CAGAGTCATTCAGAGCAATAC			
	IIICHEUKIII U	Reverse	CTTTCAAGATGAGTTGGATGG			