

Pleurotus abieticola Polysaccharide Alleviates Hyper-lipidemia Symptoms via Inhibition of Nuclear Factor- κ B/Signal Transducer and Activator of Transcription 3-Mediated Inflammatory Responses

Yongfeng Zhang¹, Yingjie Lin¹, Keyi Wu¹ Ming Jiang², Lanzhou Li^{1*}, Yang Liu^{1*}

¹ *Engineering Research Center of Chinese Ministry of Education for Edible and Medicinal Fungi, Jilin Agricultural University, Changchun, 130118, China*

² *College of Life Science and Technology, Mudanjiang Normal University, Mudanjiang, 157011, China.*

*Corresponding authors:

Yang Liu, Ph.D.

Engineering Research Center of Chinese Ministry of Education for Edible and Medicinal Fungi, Jilin Agricultural University,
Changchun, Jilin, P.R.China.

Email: y_liu10@jlau.edu.cn

Lanzhou Li, Ph.D.

Engineering Research Center of Chinese Ministry of Education for Edible and Medicinal Fungi, Jilin Agricultural University,
Changchun, Jilin, P.R.China.

Email: lilanzhou@jlau.edu.cn

Yongfeng Zhang, Email: zhangyongfeng@jlau.edu.cn

Yingjie Lin, Email: linyingjie@uor.edu.cn

Keyi Wu, Email: wukeyi@mails.jlau.edu.cn

Ming Jiang, Email: jiangming@mdjnu.edu.cn

Supplementary data

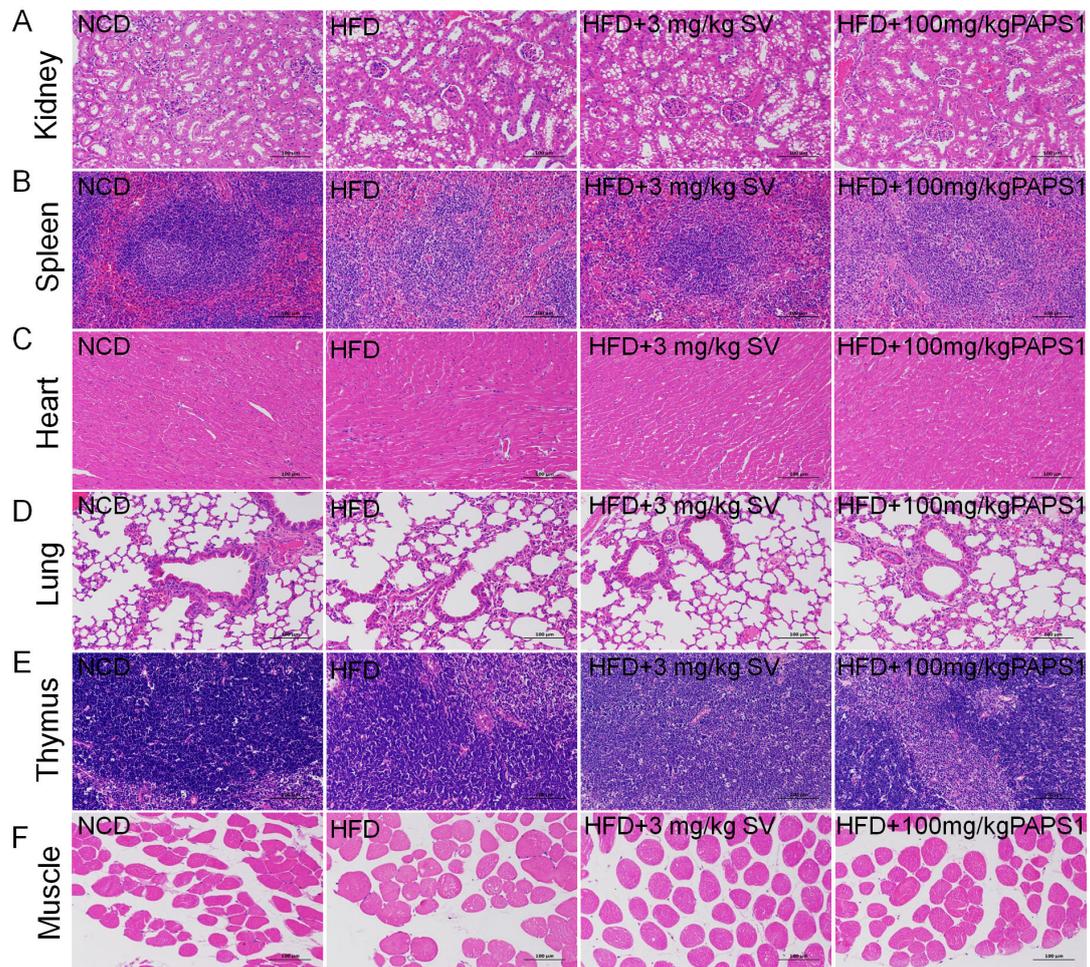


Figure S1. Histopathological analysis of the organs (kidney, spleen, heart, lung, thymus, and muscle) of mice using H&E staining (200 \times ; scale bar: 100 μ m). NCD: normal chow diet; HFD: high-fat diet; SV: simvastatin; PAPS1: *Pleurotus abieticola* polysaccharide.

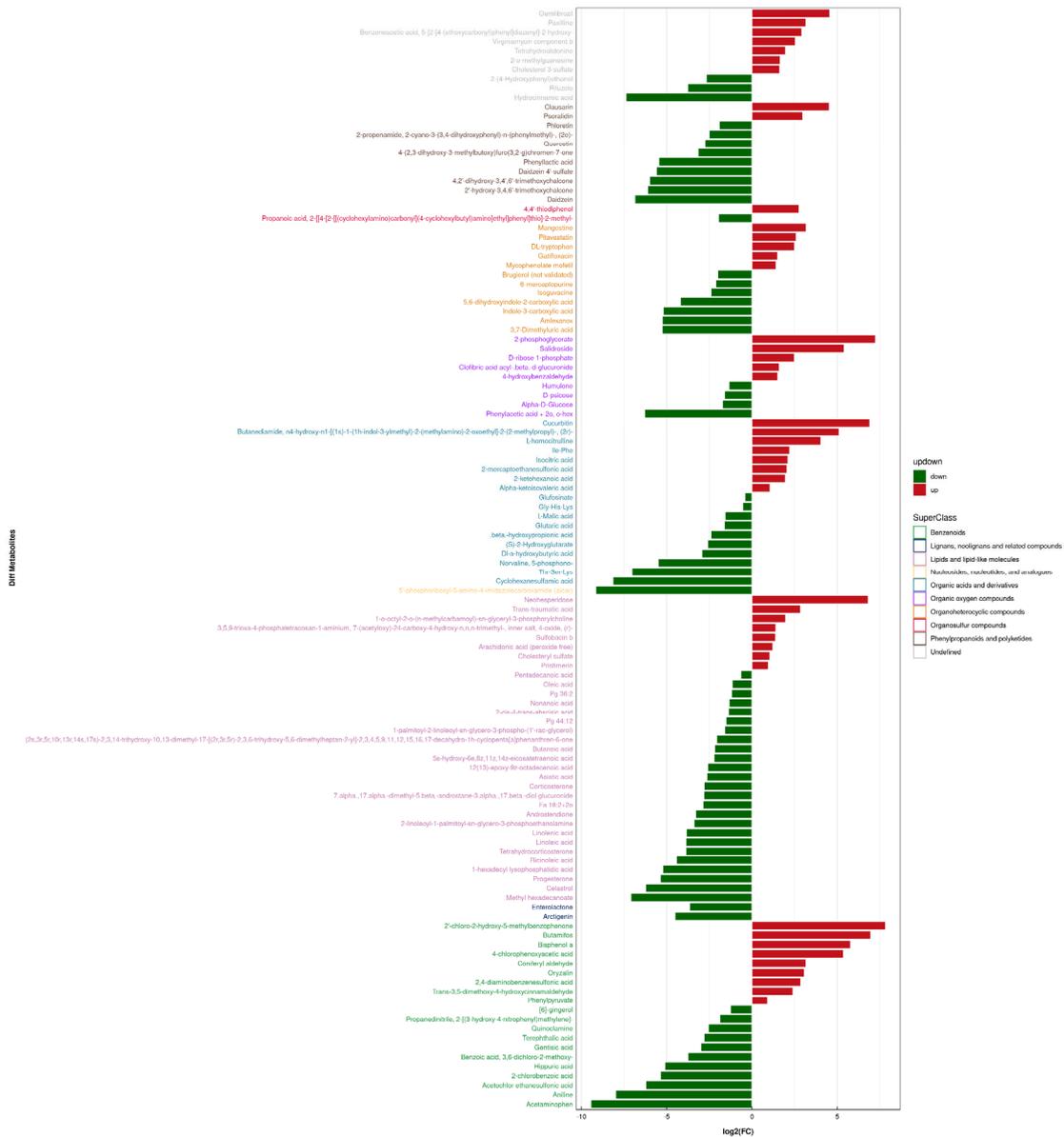


Figure S2. Differential fold analysis of the significantly differential fecal metabolites in the negative ion modes between NCD and HFD group.

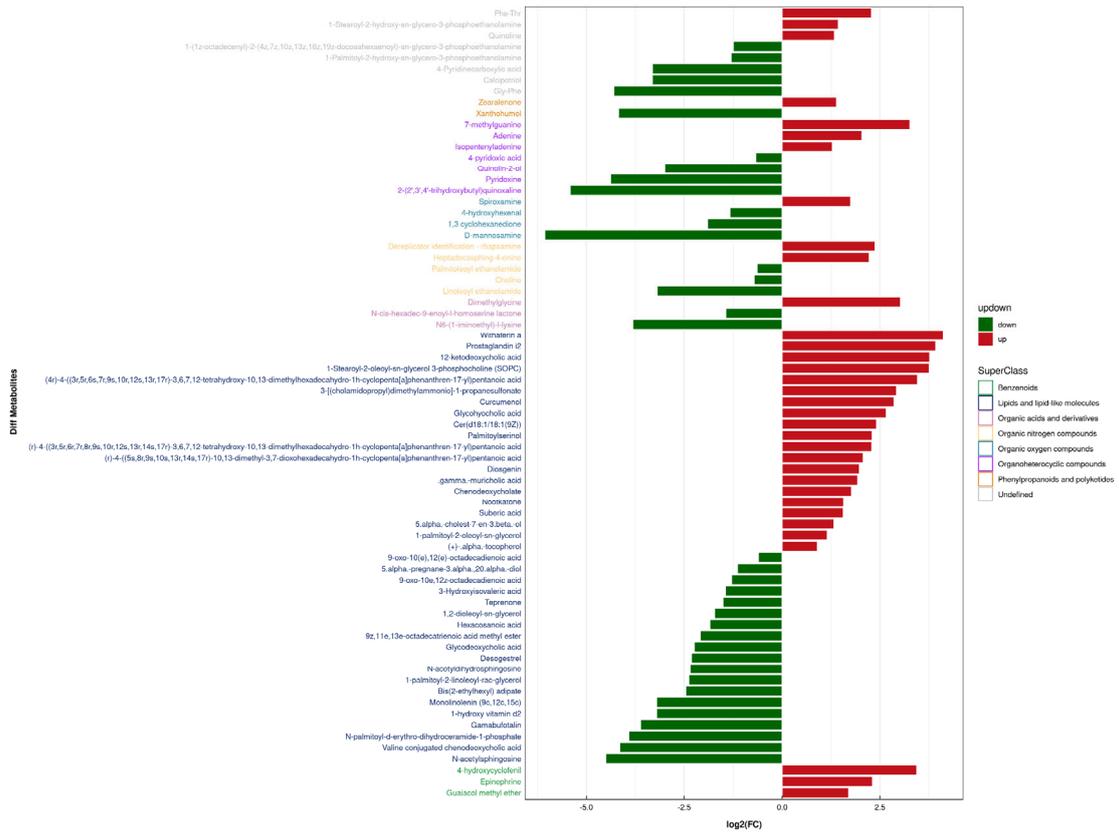


Figure S3. Differential fold analysis of the significantly differential fecal metabolites in the positive ion modes between NCD and HFD group.

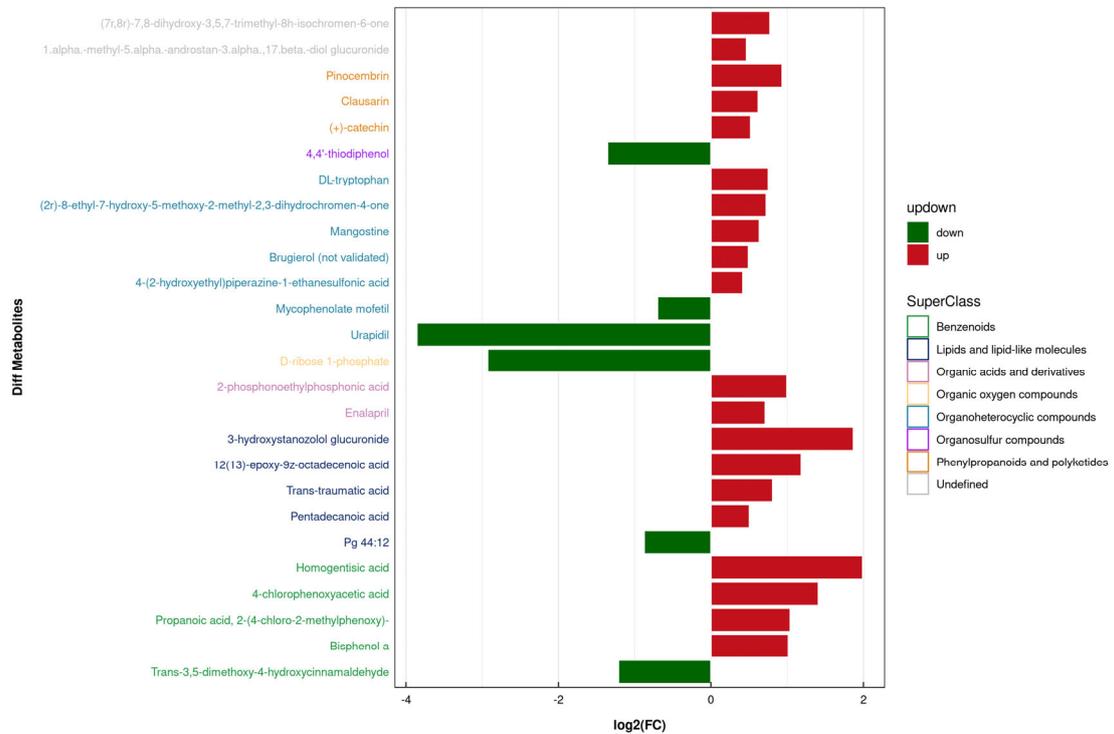


Figure S4. Differential fold analysis of significantly different metabolites in the negative ion modes between HFD and PAPS1 group.

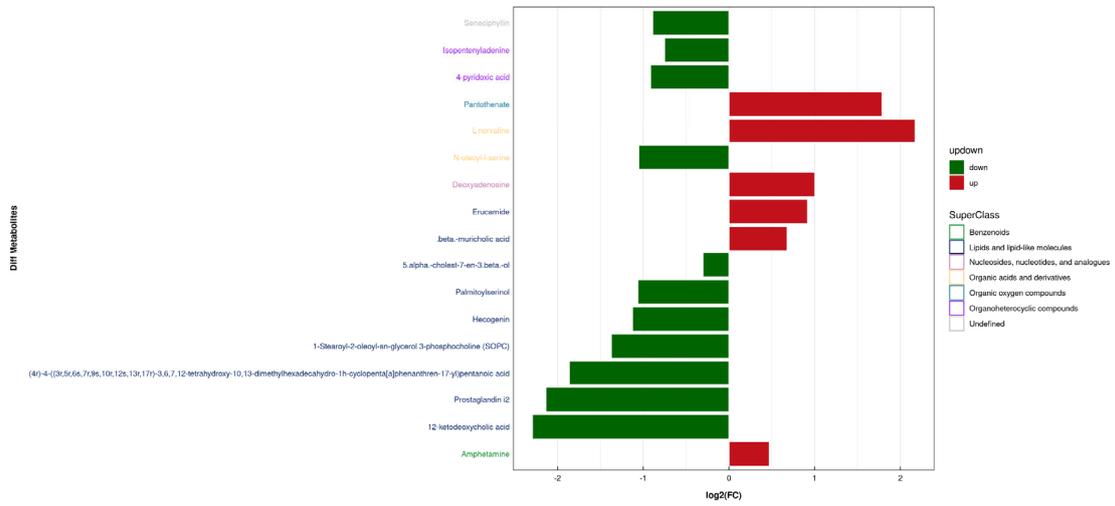


Figure S5. Differential fold analysis of significantly different metabolites in the positive ion modes between HFD and PAPS1 group.

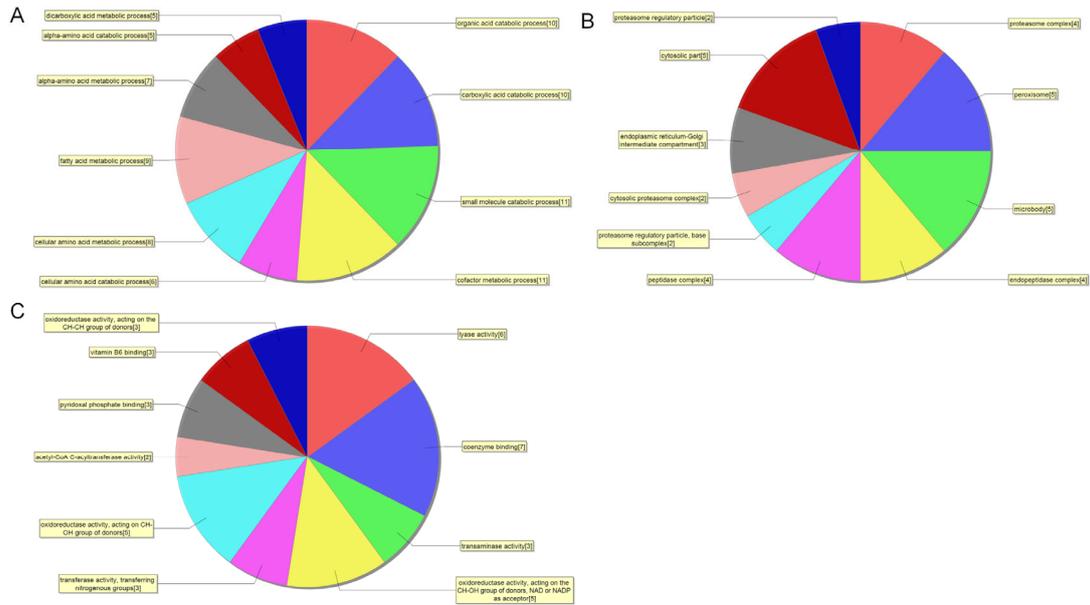


Figure S6. Pie chart of biological process (A), cellular component (B), molecular function (C) classification of GO enrichment analysis.

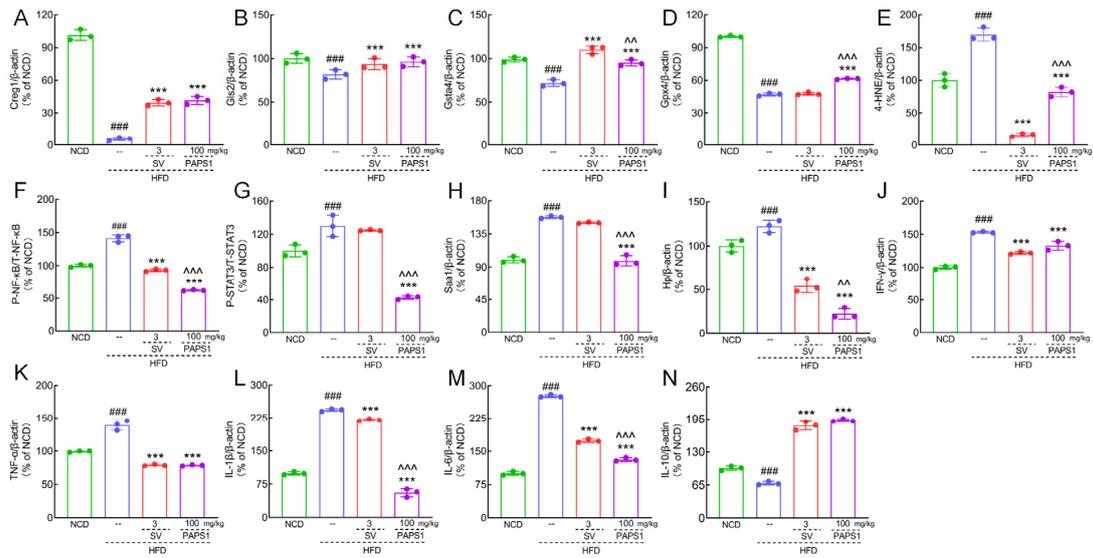


Figure S7. Quantification of protein expression in Figure 5D and 6A normalized to that of β -actin and expressed as the percentage of NCD group ($n = 3$). The data are shown as the mean \pm S.E.M. ### $p < 0.001$ versus NCD group; *** $p < 0.001$ versus HFD group; ^^ $p < 0.01$, and ^^ $p < 0.001$ versus SV group. NCD: normal chow diet; HFD: high-fat diet; SV: simvastatin; PAPS1: *Pleurotus abieticola* polysaccharide; Creg1: cellular repressor of E1A-stimulated genes 1; Gls2: glutaminase 2; Gsta4: glutathione S-transferase alpha 4; Gpx4: glutathione peroxidase 4; 4-HNE: 4-hydroxynonenal.; P-NF- κ B: phosphorylated NF- κ B; P-STAT3: phosphorylated STAT3; Saa1: serum amyloid A protein; Hp: haptoglobin; IFN- γ : interferon- γ ; TNF- α : tumor necrosis factor- α ; IL-1 β : interleukin-1 β .

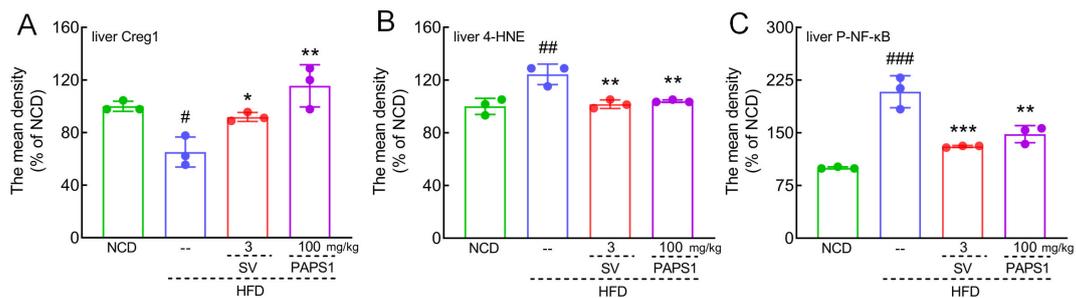


Figure S8. Quantification of protein expression in Figure 5E-G and 6I normalized to that of β -actin and expressed as the percentage of NCD mice ($n = 3$). The data are shown as the mean \pm S.E.M. # $p < 0.05$, ## $p < 0.01$ and ### $p < 0.001$ versus NCD group; * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$ versus HFD group. NCD: normal chow diet; HFD: high-fat diet; SV: simvastatin; PAPS1: *Pleurotus abieticola* polysaccharide; Creg1: cellular repressor of E1A-stimulated genes 1; 4-HNE: 4-hydroxynonenal; P-NF- κ B: phosphorylated NF- κ B.

Table S1. Details of kits used in biochemical assay and ELISA.

| Kits | Catalog number | Company | Area |
|--|----------------|--|-----------------|
| TC detection kit | A111-1-1 | Nanjing Jiancheng Bioengineering Institute | Nanjing, China |
| TC detection kit | A110-1-1 | Nanjing Jiancheng Bioengineering Institute | Nanjing, China |
| LDL-C detection kit | A113-1-1 | Nanjing Jiancheng Bioengineering Institute | Nanjing, China |
| HDL-C detection kit | A112-1-1 | Nanjing Jiancheng Bioengineering Institute | Nanjing, China |
| Mouse ALT ELISA kit | MM-44625M1 | Meimian Biotechnology | Yancheng, China |
| Mouse AST ELISA kit | MM-44115M1 | Meimian Biotechnology | Yancheng, China |
| Mouse ROS ELISA kit | MM-43700M1 | Meimian Biotechnology | Yancheng, China |
| Mouse TNF- α uncoated ELISA kit | 88-7324 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-1 β uncoated ELISA kit | 88-7013 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-6 uncoated ELISA kit | 88-7064 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-10 uncoated ELISA kit | 88-7105 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-17A uncoated ELISA kit | 88-7371 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-22 uncoated ELISA kit | 88-7422 | Thermo Fisher Scientific | MA, USA |
| Mouse IL-23 uncoated ELISA kit | 88-7230 | Thermo Fisher Scientific | MA, USA |
| BCA kit | 23225 | Thermo Fisher Scientific | MA, USA |

Table S2. Details of antibodies used in IHC and WB

| Antibody | Molecular weight | Catalog number | Dilution for WB | Dilution for IHC | Company | Area |
|------------------------|------------------|----------------|-----------------|------------------|-------------|-----------------|
| P-NF- κ B | 65 kDa | AF2006 | 1:1000 | 1:500 | Affinity | Jiangsu, China |
| T-NF- κ B | 65 kDa | A18210 | 1:2000 | / | Abclonal | Wuhan, China |
| P-STAT3 | 86 kDa | AF3293 | 1:2000 | / | Affinity | Jiangsu, China |
| T-STAT3 | 92 kDa | 3096489 | 1:2000 | / | MILLIPORE | Shanghai, China |
| IL-1 β | 17/35 kDa | A16288 | 1:2000 | / | Abclonal | Wuhan, China |
| IL-6 | 23 kDa | bs-0782R | 1:2000 | / | Bioss | Beijing, China |
| TNF- α | 17/25 kDa | A0277 | 1:2000 | / | Abclonal | Wuhan, China |
| IL-10 | 20/22 kDa | A2171 | 1:1000 | / | Abclonal | Wuhan, China |
| IFN- γ | 19 kDa | DF6045 | 1:2000 | / | Affinity | Jiangsu, China |
| Creg1 | 24 kDa | DF10194 | 1:3000 | 1:500 | Affinity | Jiangsu, China |
| Gsta4 | 27 kDa | DF12203 | 1:2000 | / | Affinity | Jiangsu, China |
| Gls2 | 66 kDa | DF13386 | 1:2000 | / | Affinity | Jiangsu, China |
| Hp | 45 kDa | 381731 | 1:1000 | / | ZEN BIO | Chengdu, China |
| Saa1 | 13 kDa | ER62519 | 1:2000 | / | HuaBio | Hangzhou, China |
| Gpx4 | 17 kDa | A19544 | 1:2000 | / | Abclonal | Wuhan, China |
| 4-HNE | 72 kDa | ab46545 | 1:2000 | 1:500 | Abcam | Cambridge, UK |
| β -actin | 43 kDa | Sc-47778 | 1:2000 | / | Santa Cruz | TX, USA |
| Goat Anti-Rabbit (H+L) | / | E-AB-1003 | 1:2000 | 1:400 | Elabscience | TX, USA |
| Goat-Anti-Mouse (H+L) | / | E-AB-1001 | 1:2000 | 1:400 | Elabscience | TX, USA |

Table S3. Formula of normal chow diet and high-fat diet.

| | NCD | HFD |
|----------------------|------|------|
| Fat (kcal%) | 10% | 60% |
| Protein (kcal%) | 20% | 20% |
| Carbohydrate (kcal%) | 70% | 20% |
| Total (kcal%) | 100% | 100% |

NCD: normal chow diet; HFD: high-fat diet; SV: simvastatin; PAPS1: *Pleurotus abieticola* polysaccharide.

Table S4. The effect PAPS1 on organ indexes in HLP mice.

| Organ | NCD | HFD | | |
|----------|-------------|----------------|--------------|-----------------|
| | | | 3 mg/kg SV | 100 mg/kg PAPS1 |
| Liver | 3.929±0.139 | 4.586±0.077## | 4.119±0.143 | 4.105±0.037* |
| Kidney | 1.451±0.032 | 1.425±0.017 | 1.475±0.041 | 1.473±0.038 |
| Spleen | 0.337±0.012 | 0.333±0.024 | 0.362±0.026 | 0.336±0.018 |
| Heart | 0.703±0.018 | 0.631±0.016# | 0.699±0.005* | 0.665±0.008 |
| Lung | 0.606±0.086 | 0.58±0.017 | 0.586±0.014 | 0.572±0.015 |
| Thymus | 0.070±0.003 | 0.052±0.003 | 0.065±0.004 | 0.073±0.007* |
| Pancreas | 0.796±0.029 | 0.590±0.017### | 0.704±0.015* | 0.737±0.026** |

The data were analyzed using a one-way ANOVA and expressed as means ± S.E.M. (n = 6). #*p* < 0.05, ##*p* < 0.01 and ###*p* < 0.001 versus NCD group, **p* < 0.05 and ***p* < 0.01 versus HFD group. NCD: normal chow diet; HFD: high-fat diet; SV: simvastatin; PAPS1: *Pleurotus abieticola* polysaccharide.

Table S5. Proteins with significantly differential expression levels in proteomics

| Number | Protein. names | Unique. peptides | fc. NCD-HFD | fc. HFD-PAPS1 |
|--|--------------------------|------------------|-------------|---------------|
| Up-regulated proteins by PAPS1 (Number:39) | | | | |
| 1 | Surf4 | 2 | 39.118139 | 0.0465283 |
| 2 | Fabp4 | 4 | 7.4147893 | 0.0712626 |
| 3 | Cyp3a11;Cyp3a16;Cyp3a41a | 6 | 75.610755 | 0.1147315 |
| 4 | Oxsm | 2 | 7.0736367 | 0.1739606 |

| | | | | |
|----|---------------------------------|----|-----------|-----------|
| 5 | Gsta4 | 2 | 4.4557578 | 0.232885 |
| 6 | Scrn2 | 2 | 5.5890002 | 0.2454149 |
| 7 | Bbox1 | 2 | 3.8376455 | 0.2995994 |
| 8 | Acaal1b | 5 | 2.0669158 | 0.3618472 |
| 9 | Sigmar1 | 2 | 2.4689319 | 0.3670756 |
| 10 | Etf1 | 3 | 4.6934167 | 0.3737625 |
| 11 | Inmt | 11 | 4.2251969 | 0.3767676 |
| 12 | ORF11;0610007P14Rik | 2 | 1.8730031 | 0.3849625 |
| 13 | Grn | 2 | 7.6496016 | 0.3984767 |
| 14 | Ca3 | 16 | 2.3182793 | 0.4141417 |
| 15 | Creg1 | 2 | 5.9268832 | 0.4161873 |
| 16 | Cyp2a5;Cyp2a4 | 5 | 5.5953594 | 0.4324899 |
| 17 | Ehhadh | 30 | 1.5761045 | 0.4836167 |
| 18 | Mtstp8 | 2 | 2.6396857 | 0.498133 |
| 19 | Ddah1 | 7 | 3.7453863 | 0.5263316 |
| 20 | Mpc2 | 4 | 1.6294373 | 0.5414203 |
| 21 | Sec24d | 2 | 4.4387162 | 0.5575168 |
| 22 | Coq9 | 3 | 1.7423806 | 0.5597148 |
| 23 | Oat | 11 | 1.9601638 | 0.5867929 |
| 24 | Usp14 | 2 | 2.480496 | 0.5873441 |
| 25 | Acaal1a | 4 | 1.8526392 | 0.593132 |
| 26 | Khk | 11 | 2.9074653 | 0.6037921 |
| 27 | Ces1f | 15 | 2.658276 | 0.6038403 |
| 28 | Me1 | 13 | 3.0997432 | 0.6125076 |
| 29 | Acad8 | 8 | 1.6742251 | 0.6209792 |
| 30 | Bdh1 | 9 | 1.598835 | 0.6212225 |
| 31 | Gck | 5 | 1.57656 | 0.6235898 |
| 32 | Hibadh | 8 | 1.5479204 | 0.6323804 |
| 33 | Cisd1 | 4 | 2.2395791 | 0.6401977 |
| 34 | Cyp2c67;Cyp2c68;Cyp2c69;Cyp2c40 | 7 | 2.1551439 | 0.6405674 |
| 35 | Gstt1 | 8 | 1.6869598 | 0.6471396 |
| 36 | Hal | 14 | 2.5095091 | 0.6494651 |
| 37 | Pecr | 5 | 1.5320279 | 0.6512129 |
| 38 | Gls2 | 8 | 2.7497557 | 0.6537813 |
| 39 | Gpt2 | 5 | 2.0618183 | 0.6592659 |

Down-regulated proteins by PAPS1 (Number:25)

| | | | | |
|---|--------|---|-----------|-----------|
| 1 | Ptms | 2 | 0.2024234 | 1.5057212 |
| 2 | Kng1 | 3 | 0.6039607 | 1.5063507 |
| 3 | Mbl1 | 2 | 0.1350671 | 1.5108965 |
| 4 | Hnrnpm | 5 | 0.2355895 | 1.5144546 |
| 5 | Lrpap1 | 2 | 0.6098233 | 1.5453759 |

| | | | | |
|----|-----------|----|-----------|-----------|
| 6 | Creld2 | 3 | 0.1830977 | 1.5608068 |
| 7 | Rer1 | 2 | 0.3462425 | 1.5711873 |
| 8 | Rpl13a | 5 | 0.5687837 | 1.5767496 |
| 9 | Slc37a4 | 2 | 0.5201448 | 1.6010325 |
| 10 | Psemb7 | 2 | 0.2572866 | 1.619552 |
| 11 | Psmc2 | 3 | 0.2652712 | 1.6283054 |
| 12 | Rdh11 | 2 | 0.655161 | 1.6459288 |
| 13 | Rps28 | 2 | 0.1257974 | 1.6765039 |
| 14 | Canx | 15 | 0.5321497 | 1.734502 |
| 15 | Rmdn3 | 3 | 0.1884984 | 1.735627 |
| 16 | Apex1 | 2 | 0.4296374 | 1.7507134 |
| 17 | Rps6 | 5 | 0.317078 | 1.7868004 |
| 18 | Psmd2 | 8 | 0.6314325 | 1.8031546 |
| 19 | D10Jhu81e | 7 | 0.5121806 | 1.8408467 |
| 20 | Ccbl2 | 6 | 0.6094626 | 1.8646155 |
| 21 | Acbd5 | 3 | 0.3950492 | 2.4316103 |
| 22 | Hp | 11 | 0.546026 | 3.1897325 |
| 23 | Saa1 | 2 | 0.1559159 | 4.0626199 |
| 24 | Tagln2 | 2 | 0.2171844 | 4.3434152 |
| 25 | Rps15 | 2 | 0.3475577 | 4.4990631 |

fc. NCD-HFD: the ratio of protein between NCD group and HFD group;

fc. HFD-PAPS1: the ratio of protein between HFD group and PAPS1 group.

NCD: normal chow diet; HFD: high-fat diet; PAPS1: *Pleurotus abieticola* polysaccharide.