

# Function and structure of a novel anti-diabetes agent from Ganoderma Lucidum

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# INTRODUCTION



Morris F White. Insulin signaling in health and disease Science, 2003, 302 (5651):1710-1711

 Inhibition of protein tyrosine phosphatase 1B (PTP1B) activity has been considered as a promising therapy approach to treat type 2 diabetes.

• Protein tyrosine phosphatase 1B (PTP1B) have been implicated in the regulation of insulin signal transduction process

• PTP1B dephosphorylate the insulin receptor as well as the substrate proteins, controlling the insulin signaling pathway

• Overactivation of PTP1B inhibits the insulin receptor signaling cascade. Therefore, PTP1B is an insulinsensitive drug target for anti-diabetes.

## OBJECTIVE

In this work, a novel PTP1B activity inhibitor, named FYGL (Fudan-Yueyang-G. lucidum), was screened from the fruiting bodies of Ganoderma lucidum. The efficient PTP1B inhibitory potency, plasma glucose level in vivo, toxicity of FYGL, and structure of FYGL were studied.

# RESULTS



Bao-Song Teng, Ping Zhou,\* et al. J. Agric. Food Chem. 2011, 59(12), 6492-6500.

Ping Zhou



Normal Control 75mg/kg 250mg/kg 450mg/kg 200mg/kg FYGL FYGL FYGL Mefformin

n = 8, \*p < 0.05 vs. control, \*\*p < 0.01 vs. control.

### • HbAlc is considered a "golden index" indicating the plasma glucose level. After 8 weeks, HbA1c level was significantly decreased dose-dependently for the mice treated by FYGL and metformin.

Deng Pan, Ping Zhou,\* et al. *PLoS One*, 2013, 8(7), e68332.









metformin 200mg/kg

 $6.4 \pm 0.3$  \*\*

PTP1B activity in skeletal muscle

Compared with control group, PTP1B expression and activity were inhibited dose-dependently in FYGL group, also indicating that the target of FYGL is PTP1B in vivo. Chendong Wang, Ping Zhou,\* et al. Brit J Nutr, 2012, 108, 2014-2025.

be represent by the following groups:  $\rightarrow 2.4$ )- $\alpha$ -L-Rhap-(1-

R→2)-α-D-Glcp-(3-

 $\beta$ -D-Galp-(1 $\rightarrow$ 3)-Ga

 $\rightarrow 2$ )- $\alpha$ -L-Rhap-(1 $\rightarrow 2$ )- $\alpha$ 

protein-T





# RESULTS

### Structure characteristic of FYGL



• Figure is  $\beta$ -elimination reaction probed by UV, which indicates that protein is bound with saccharide by O-glycosidic linkage.

• Table is amino acid contents before and after  $\beta$ -elimination reaction, which shows that after  $\beta$ elimination reaction, both Thr and Ser contents were decreased, while Ala increased, indicating that protein bind covalently with saccharide by Thr and Ser residues.



• NMR analysis suggest *FYGL* being a heteropolysaccharide with  $\alpha$  and  $\beta$  linkages, the peaks within 170 - 175 ppm in <sup>13</sup>C NMR indicates protein present. The backbone is hyperbranced polysaccharide and proteins are grafted. (Deng Pan, Ping Zhou, \* et al. Carbohydrate Polymer, 2015, 117, 106–114.)

## CONCLUSION

- FYGL, screened from G. lucidum, is an efficient PTP1B inhibitor in vivo
- FYGL can decrease the plasma glucose level through inhibiting the PTP1B expression and activity, consequently, regulating the tyrosine phosphorylation level of the IR β-subunit.
- 3. FYGL contain hyperbranched proteoglycan, which may play special roles for its bioactivities of PTP1B inhibition and antihyperglycemic potency.

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α-D-Glcp

 $\rightarrow$ 6)-B-D-Galp-1 $\rightarrow$ , Araf-(1 $\rightarrow$  and  $\rightarrow$ 3.6)-B-D-Galp-(1

R→2)-α-D-Glci