Whole-genome identification and expression pattern of the vicinal oxygen chelate family in Rapeseeed (*Brassica napus* L.)

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Supplementary Information:

Image legends

Image S2. Predicted secondary structure of BnaGLYI. BnaGLYII was an example to show that results from GOR4 prediction and PSIPRED is relatively the same. A: Results from GOR4 prediction analysis. B: Results from PSIPRED prediction analysis.

Image S2

A 10 20 30 40 50 60 70

Sequence length: 274

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GOR4:
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(Hh) : 47 is 17.15% Alpha helix 3₁₀ helix (Gg) : 0 is 0.00% Pi helix (Ii) : 0 is 0.00% Beta bridge (Bb) : 0 is 0.00% Extended strand (Ee) : 78 is 28.47% (Tt): 0.00% Beta turn 0 is Bend region (Ss) : 0.00% 0 is (Cc) : 149 is 54.38% Random coil Ambiguous states (?) 0 is 0.00% Other states 0 is 0.00%

