

Coprophagous features in carnivorous *Nepenthes* plants: a task for ureases

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Supplementary Information:

Figure S1. Homology among amino acid sequences of ureases. Included are ureases from seven carnivorous plants (*Aldrovanda vesiculosa*, *Cephalotus follicularis*, *Dionaea muscipula*, *Drosera spatulata*, *Genlisea aurea*, *Nepenthes alata*, *Nepenthes hemsleyana*) and three non-carnivorous plants (*Canavalia ensiformis*, *Glycine max* embryo-specific, *Arabidopsis thaliana*). Different colors indicate individual amino acids that are different to the consensus amino acid at a particular position.

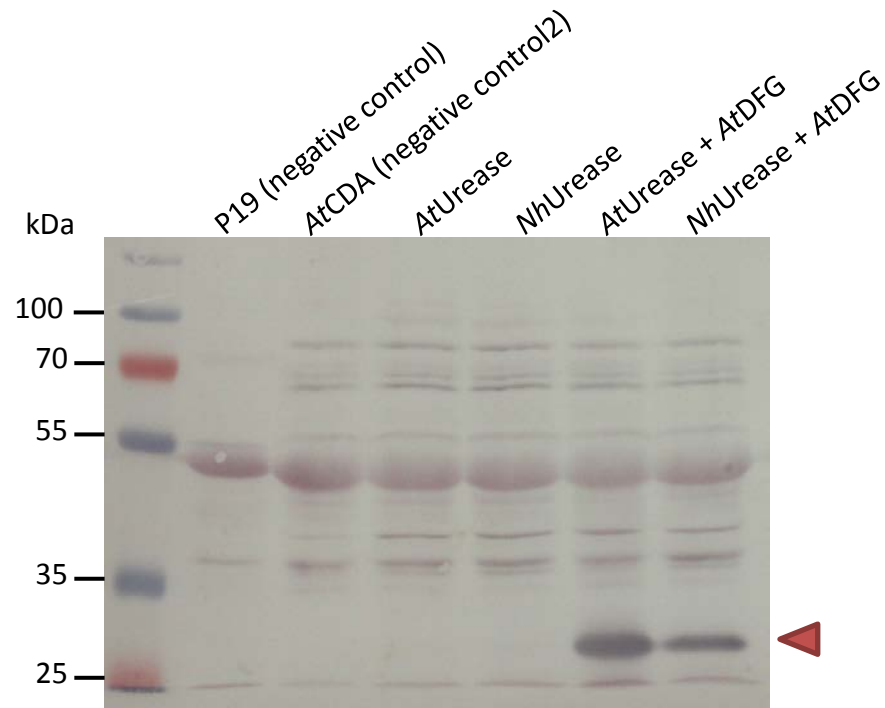


Figure S2. Western blot of *Arabidopsis thaliana* AtUreD heterologously co-expressed with urease in *Nicotiana benthamiana*. Full size membrane of the Western blot. 10 μ l of each sample was loaded to a 10% SDS gel. The samples are the same as used in Figure 3; additionally with one more negative control of a heterologously expressed CYTIDINE DEAMINASE (*AtCDA*) (Ref: Chen et al., 2016, Plant Physiol. 171, 799-809; DOI: 10.1104/pp.15.02031). All lanes were analysed by immunoblot employing specific AtUreD antiserum. The triangle indicates the targeted protein AtUreD.

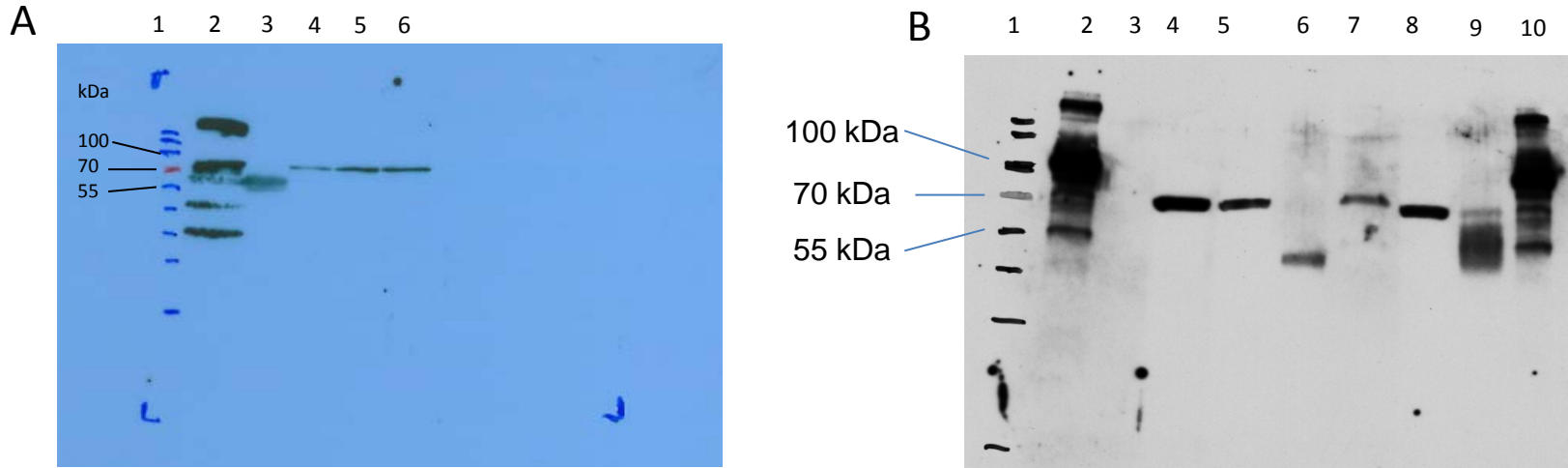


Figure S3. Western blot analyses of urease proteins. Occurrence of urease was done in crude protein extracts from *Nepenthes alata* and *N. hemsleyana* by immunoblot using polyclonal anti-jackbean urease antibodies. (A) Lanes: 1. Page ruler prestained protein ladder (Thermo Fisher); 2. 5 μ g total protein extract from soybean seeds; 3. 1 μ g of 6 months old degraded total protein extract from *N. alata* pitcher; 4. 1 μ g from *N. alata* leaf; 5. 1 μ g total protein extract from *N. alata* pitcher; 6. 1 μ g total protein extract from another *N. alata* pitcher; Lanes 4, 5, 6 are shown in Figure 2A. (B) Lanes: 1. Page ruler prestained protein ladder; 2. 200 ng urease from *Canavalia ensiformis* (jackbean) as positive control (Sigma-Aldrich); 3. Empty lane; 4. 5 μ g total protein extract from *N. hemsleyana* leaf; 5. 5 μ g total protein extract from *N. hemsleyana* pitcher; 6. 5 μ g, 6 months old degraded total protein extract from *N. hemsleyana* pitcher; 7. 5 μ g total protein extract from *N. alata* leaf; 8. 5 μ g total protein extract from *N. alata* pitcher; 9. 5 μ g, 6 months old degraded total protein extract from *N. alata* pitcher; 10. 200 ng urease from *C. ensiformis*. Lanes 4, 5 are shown in Figure 2A.

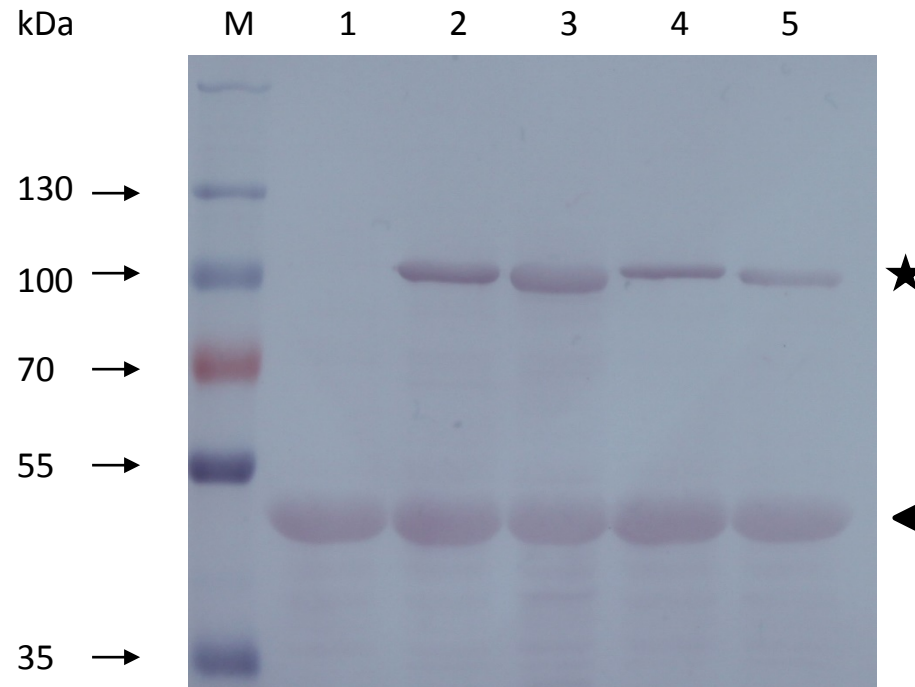


Figure S4. Western blot of *Nepenthes hemsleyana* urease heterologously expressed in *Nicotiana benthamiana*. The full size membrane of the Western blot in Figure 3. Lane 1 to 5 represent extracts of transiently expressed P19: Lane 1 negative control; 2: *A. thaliana* urease; 3: *N. hemsleyana* urease; 4: *A. thaliana* urease with accessory proteins UreD, UreF and UreG; 5: *N. hemsleyana* urease with accessory proteins UreD, UreF and UreG. Western blot was developed with anti *A. thaliana* urease antiserum. The star indicates the targeted protein urease and the triangle indicates Rubisco of *N. benthamiana*.