Functional characterization of a class III acid endochitinase from the traps of the carnivorous pitcher plant genus, *Nepenthes*

Sandy Rottloff, Regina Stieber, Heiko Maischak, Florian G Turini, Günther Heubl, and Axel Mithofer

Fig. S1. Amino acid sequence alignment of endochitinases from various species. The amino acid sequence alignment and the deduced consensus sequence was generated using Clustal X involving the plant species *Pyrus pyrifolia; Lupinus albus; Cucumis sativus; Nepenthes rafflesiana; Beta vulgaris; Vitis vinifera; Oryza sativa.*

P.pyrifolia L.albus C.sativus N.rafflesiana B.vulgaris V.vinifera O.sativa consensus	1 MASKTQTLALTLSLLILISS-CKSSQAAG-HAIYWGONGNEGTLADACNSGNYQFVNIAELITEGNNOTEVLNLAGHCDPASGTCTGLSA 1 MASLKQVSLILFPLLLLISSSFKLSNAAG-IVIYWGONGNEGSLADACNTNNYQYVNIAELSTFGNGOTPELNLAGHSRDGLNA 1 MAAHKITTTLSIFFLLSSIFRSSDAAG-IAIYWGONGNEGSLASTCATGNYEFVNIAELSSFGSGOAPVLNLAGHCNPDNNGCAFLSD 1 MATHYSSAILPILTLFVFLS-INPSHGSG-IAYWGONGNEGTLSDTCATGNYNYVLVSELTTEGNGOTPVLNLAGHCDPSSNGCTGLST 1 MAAKIVSVLFLISLLIFASFESSHGSQ-IVIYWGONGDEGSLADTCNSGNYGTVILAEVATFGNGOTPVLNLAGHCDPASNGCTGSS 1 MAAKIVSVLFLISLLIFASFESSHGSQ-IVIYWGONGDEGSLADTCNSGNYGTVILAEVATFGNGOTPLNLAGHCDPATN-CNSLSS 1 MARTPQSTPLLISLSVLALLQTSYAGG-IAIYWGONGNEGTLTQTCNTGKYSYVNIAELNKFGNGOTPEINLAGHCDPASNGCTSVST 1 MTSRMFSAMQMLIMVVVALAGLAAGTRAGDIAIYWGONGNEGTLAQTGATGNYRFYIVAELPVFGKGOTPVLNLAGHCDPASNGCTGVGA 4 *
P.pyrifolia L.albus C.sativus N.rafflesiana B.vulgaris V.vinifera O.sativa consensus	89 DIRTCOSKNIKVILSIGGASCSYSITSADDAROVADYIWNNFLGGOSASRELCDAVLDGVDFDIEACGGOOFYDELARSINGHAGOAK 84 DIKGCOGKGIKVILSIGGACSYSINSADDATNLANYIWNNFLGGTSDSRPFGDAVLDGIDFDIEAGGAOHYDELARAINGFSSOKK 88 EINSCKSONVKVILSIGGACSYSISSADDAKOVANFIWNSYLGGOSDSRPLGAAVLDGVDFDIESCSGOFWDVLAOELKNFG 89 DITSCONOGIKVILSIGGACSYSISSADDAKOVANFIWNSYLGGOSDSRPLGAAVLDGVDFDIESCSGOFWDVLAOELKNFG 87 DIKTCOOAGIKVILSIGGACGYSISSADDAKOVANFIWNYLGGOSDSRPLGAAVLDGIDFDIESCSOFWDVLAOELKNFG
P.pyrifolia	176 TVYLAAAPQCPIPDAHLDGAIQTGLEDYVWVQEYNNPPGQYADG-NANAFLNSWSQWAS-VPATQVFMGLPASTDAAGSG-FIPADALKS
L.albus	171 -VYLGAAPQCPIPDAHLDAAINTGLEDYVWVQEYNNPGQYACG-NTNNFINSWNQWTS-SQAKQVFLGLPASEAAAPSGGFIPTDVLIS
C.sativus	171 QVILSAAPQCPIPDAHLDAAIKTGLEDSVWVQEYNNPPGMFADNADNLLSSWNQWTA-FPTSKLYMGLPASEAAAPSGGFIPTDVLIS
N.rafflesiana	173 SVLVSAAPQCPYPDAHLDLAIATGIEDYVWVQEYNNEQGEYVTDDTNLLSSWNQWTS-SQANVVFLGLPASTDAASSG-YISPDVLIS
B.vulgaris	174 TVYLSAAPQCPLPDASLSTAIATGLEDYVWVQEYNNEQGEYVTSADNLLSSWNQWTS-SQANVVFLGLPASTDAASSG-YISPDVLIS
V.vinifera	178 KVYLTAAPQCPLPDASLSTAIATGLEDYVWVQFYNNPPGQYSSG-NTNNLLSSWNQWTT-VQANQIFLGLPASSAAAGSG-FIPADALTS
O.sativa	181 PVYLTAAPQCPFPDASLGVALSTGLEDYVWVQFYNNPPGQYSSG-NTNNLLNSWNRWTSSINSTGSFMGLPASSAAAGSG-FIPADAUTS
consensus	181 ******
P.pyrifolia	263 QVLPTIKNSAKYGGVMLWSRWYDINSGYSASIKDSI-
L.albus	258 QVLPTIKTSPKYGGVMLWNGFNDIQTGYSDAIKASV-
C.sativus	259 QVLPTIKASSNYGGVMLWSKAFDNGYSDSIKGSIG
N.rafflesiana	259 QVLPSIKASSKYGGVMLWSKYYDNGYSSAIKDSV-
B.vulgaris	260 QVLPTIKGSAKYGGVMLWSKAYDSGYSSAIKSSV-
V.vinifera	266 QILPVIKRSPKYGGVMLWSKYYDDQSGYSSSIKSSV-
O.sativa	269 KVLPVVKKSPKYGGIMLWSRYYDGLTGYSDKVKSSV-
consensus	271 ** * * ***

Fig. S1