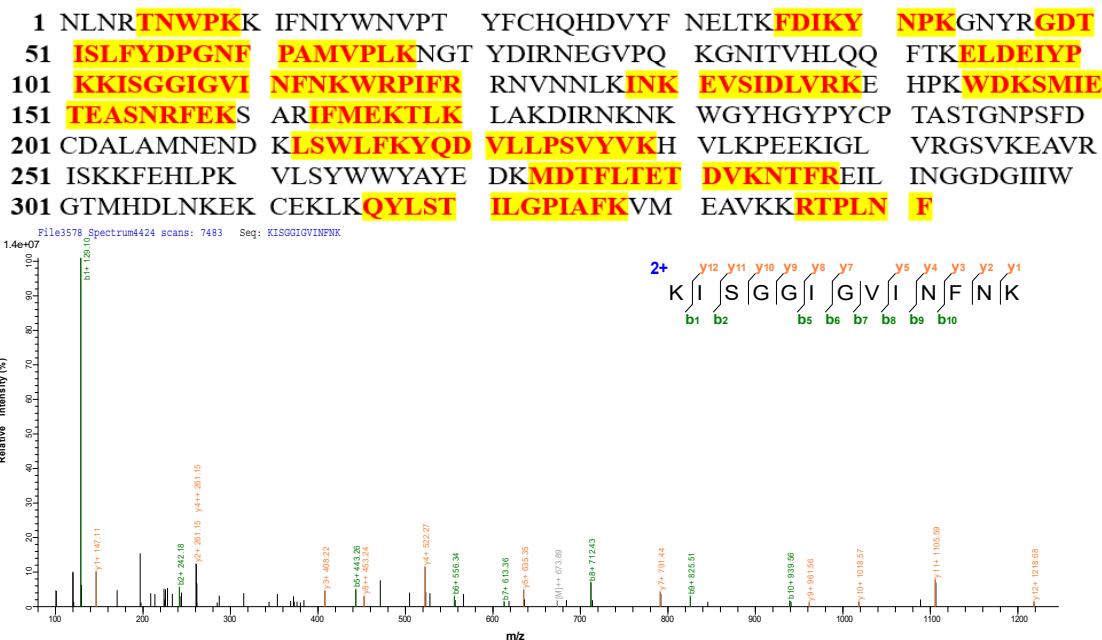


Supplementary Materials: Characterization of the Composition and Biological Activity of the Venom from *Vespa bicolor* Fabricius, a Wasp from South China

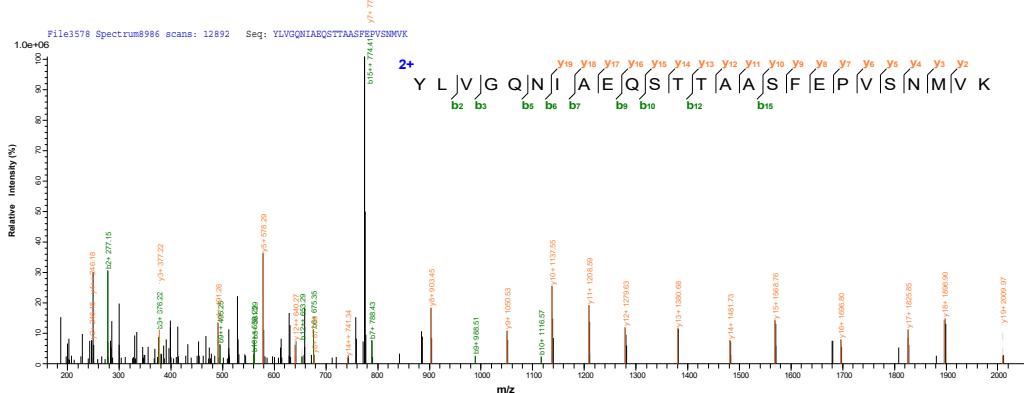
Yonghua Wu, Yu Zhang, Danqiao Fang, Jing Chen, Jingan Wang, Lin Jiang and Zhufen Lv

A. Hyaluronidase A



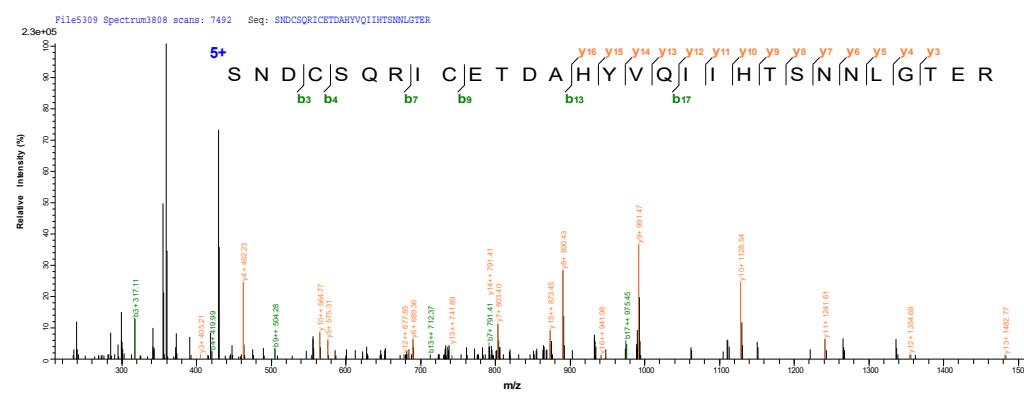
B. Allergen 5

1 NNYCKIKCRS GIHTLCKYGT STKPNCGR**SV** **VKASGLTKAE** **KLEILK**QHNE
 51 FRQKV**AR****GLE** **TRGNPGPQP**P **AKSMNTLVWN** DELAQIAQVW ASQCKYGHDN
 101 CRNTAK**YLVG** **QNIAEQSTTA** **ASFEPVS**NMV **KMWSDEVKDY** **QYGSSK**NK**LN**
 151 **DVGHYTQM**VW **AK**TKEIGCGN IKYIENGWHH HYLVCNYGPA GNIGNEPIYE
 201 KK



C. Phospholipase A1

1 GLLPKCK**LVP** **EQISFILSTR** **ENRNGVFLTL** **DSLKK**GGILN KSDLSSSTQVV
 51 FLIHGFISSA NNSNYMDMTK ALLEKNDCMV ISIDWRNGAC TNEFQILK**FI**
 101 **GYPKAVENTR** **TVGKYIADFS** **KLLMQKYKVS** **LANIRLIGH**S **LGAQIAGFAG**
 151 **KEYQKFKLGK** **YPEIIGLDPA** **GPLFKSNDCS** **QRICETDAHY** **VQIHTSNNL**
 201 **GTER**TLGTVD FYMNNGYNQP GCYYSFIGET CSHTR**AVQYF** **TECIRHECCL**
 251 **IGVPQSKNPQ** **PVKCTRNEC** **VCVGLNAKRY** **PKTGSFYVVP** **ESKAPYCNNK**
 301 GKKI



D. Phospholipase A1

1 **FNPCPYSDDT VKMILTR**EN K**KHDFYTLDT IKKHNEFK**KS TIKHQVVFIT
 51 HGFSSADTE NFLAMAKALS DKGNYLVILI DWRVAACTEE MSGIQLAYYS
 101 YAASNTR**LVG NYIATVTKML VQKYNVPMAN IRLIGHSLGA HTSGFAGKKV**
 151 **QELGLGKYSE IIGLDPAGPS FKSNDCSERI CKTDAHYVQI IHTSNHLGTL**
 201 VTLGTVDFMN NGYNQPGCGL PLIGETCSHT RAVK**YFTECI KHECCLIGVP**
 251 **QSKKPQPVSK CTRNECVCGV LNAKTYPKTG SFYVPVESKA PYCNNKGKII**

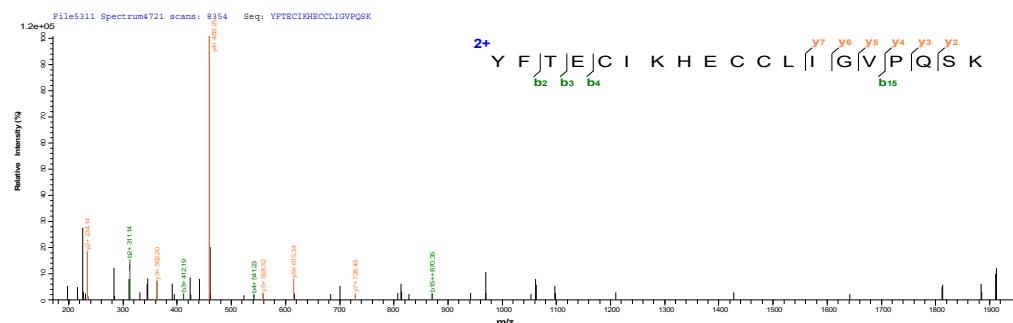
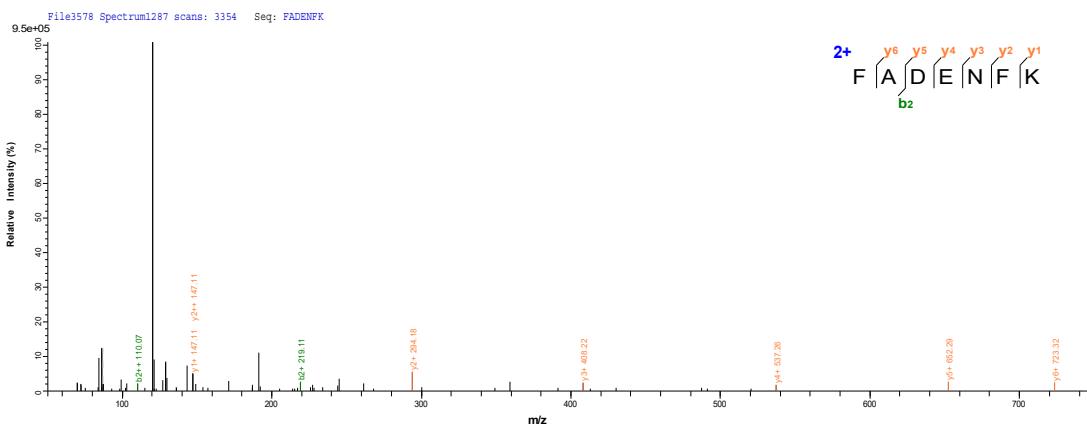


Figure S1. Four proteins in VBV were identified by LC-MS/MS (A–D), and the respective secondary mass spectrograms of proteins 1 to 4 are displayed below. A) Complete coding sequence of hyaluronidase A is shown, and a sequence coverage of the matched peptides of protein 1 in hyaluronidase A is highlighted. B) Complete coding sequence of allergen 5 is shown, and a sequence coverage of the matched peptides of protein 4 in allergen 5 is highlighted. C) Complete coding sequence of phospholipase A1 is shown, and a sequence coverage of the matched peptides of protein 2 in phospholipase A1 is highlighted. D) Complete coding sequence of phospholipase A1 protein is shown, and a sequence coverage of the matched peptides of protein 3 in phospholipase A1 is highlighted.

A. Peptidyl-prolyl cis-trans isomerase

1 MKLLLFLGLV AAASCANNAN GPKVTDKVWF DIKIGDADAG RIEIGLFGKT
 51 VPKTVKNFVE LAKKPEGEGY KGSVFHRVIK DFMIQGGDFT KGDTGGRSI
 101 YGDRFADENF KLKHYGAGWL SMANAGKDTN GSQFFITVKQ TPWLDGRHVV
 151 FGKIIKGMDI VRKIEKTNTD SRDRPQEDVV IADSGAETVA DPPSVSKDDA
 201 IN



B. Uncharacterized protein

1 MSYDLRSGMW YFDGGQPEVN LRFTRQYRLL LAPPRALPKV PRGVPSXXXX
 5 1 XXMVARALII LIINRLEQLT PAPYTKGITD LRGGSRQNRS ANNRLPLIAK
 101 QSLLRNVVIT PL

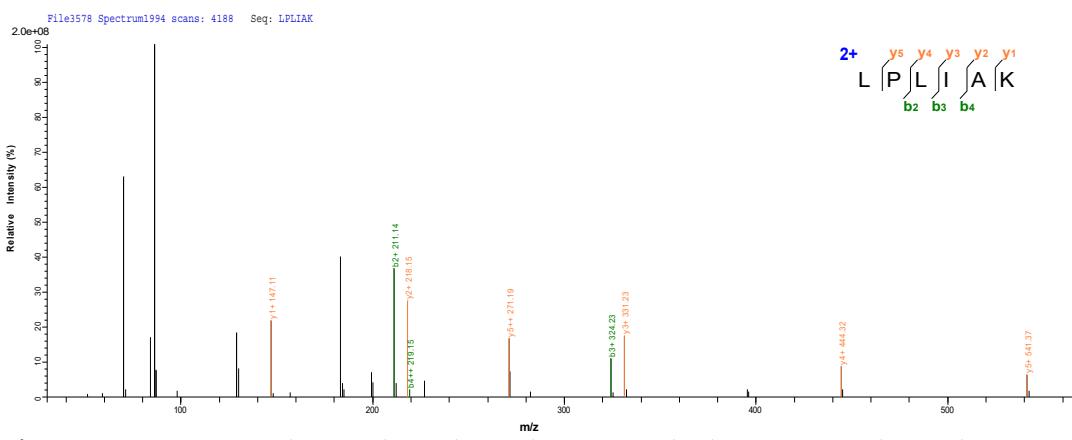


Figure S2. Proteins 5 and 6 match results in the *Apocrita* database. **A)** Complete coding sequence of peptidyl-prolyl cis-trans isomerase is shown, and a sequence coverage of the matched peptides of protein 5 in peptidyl-prolyl cis-trans isomerase is highlighted. **B)** Complete coding sequence of uncharacterized protein is shown, and a sequence coverage of the matched peptides of protein 6 in uncharacterized protein is highlighted.

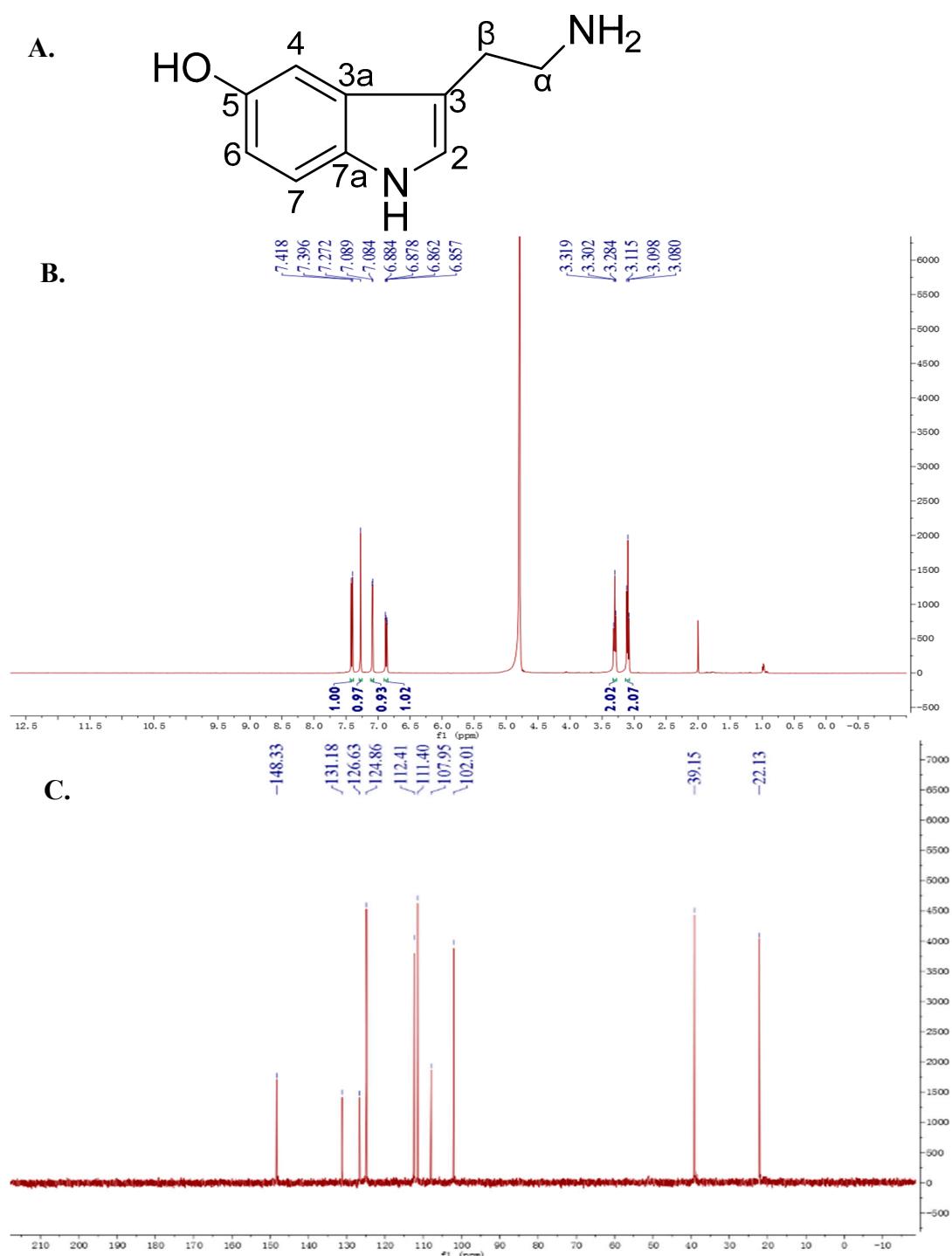


Figure S3. Component 7 of VBV was identified as 5-hydroxytryptamine (5-HT) by NMR. **A**) Structural formula of 5-HT. **B**) ^1H NMR spectrum of 5-HT in D_2O (400 MHz). **C**) ^{13}C NMR spectrum of 5-HT in D_2O (500 MHz).

Vb-MLP 12a

1 INWK^GIAMA KKLL

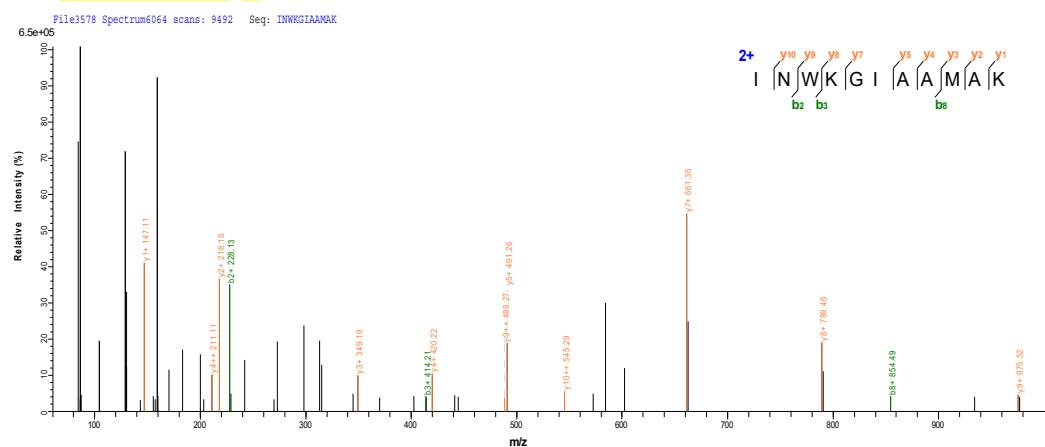


Figure S4. Component 8 of VBV was identified as Vb-MLP 12a by LC-MS/MS. The sequence coverage of the matched peptides is highlighted, and the secondary mass spectrograms of Vb-MLP 12a are displayed below.