

Table S1: Primers used in this study

| Name | 5'- 3' sequence | Purpose |
|--------------------------|---|--|
| pHbl_up1 pHbl_up1 | tgtttggttcaatTTTatggatca ttggcattgcgagtgTattg | PCR of <i>hbl-2</i> upstream region |
| pHbl_down1 pHbl_down2 | ttggaaaacattaagagcagatg tcgcaatcatagatcgtttcc | PCR of <i>hbl-2</i> downstream region |
| Hbl-LF1 Hbl-LF2 | <i>ccccgaattc</i> ggaagctttgcagctttt <i>cccactagtcc</i> gctctctttgtaaactgga | Hbl genetic deletion, upstream region |
| Hbl-RF1 Hbl-RF2 | <i>ccccgaattc</i> ccaagtaattggagctggaga <i>cccactagtac</i> aggaggtcccagtacc | Hbl genetic deletion, downstream region |
| rHblBfw rHblBrv | <i>aacat</i> atgttggaggaaaatgaaatgat <i>AAGGATCC</i> <u>tattagtgatggtgatgatgatg</u> ttttgtggagtaacagttcca | Cloning of HblB |
| rHblL1fw rHblL1rv | <i>ggatta</i> atgaaaaaattccattcaaagtactaac <i>aaggatcct</i> <u>tattagtgatggtgatgatgatg</u> ctcctgtttaaagcaatatctttg | Cloning of Hbl- L ₁ |
| rHblL2fw rHblL2rv | <i>aacat</i> atgaaaactaaaataatgacaggattt <i>aaggatcct</i> <u>tattagtgatggtgatgatgatg</u> aaattatatacttgttcttcaaggtaactta | Cloning of Hbl- L ₂ |
| rNheAfw rNheArv | <i>aacat</i> atgaaaaagactttaattacagggttattg <i>aaggatcct</i> <u>tattagtgatggtgatgatgatg</u> tacttcaacgttgtaacgtaac | Cloning of NheA |
| rNheBfw rNheBrv | <i>aacat</i> atgacaaaaaacatataaaagtaatgg <i>aaggatcct</i> <u>tattagtgatggtgatgatgatg</u> ctttttcgtgtctactactttaatc | Cloning of NheB |
| rNheCfw rNheCrv | <i>aacat</i> atgcagaaacgattttataaaaaatg <i>aaggatcct</i> <u>tattagtgatggtgatgatgatg</u> ctttgccacacctcatgtaatt | Cloning of NheC |

Italicized letters indicate restriction recognition sites for *Bam*HI, *Ase*I, *Nde*I, *Eco*RI, or *Spe*I. Underlined letters indicate histidine tags and stop codons.