

## Corrigendum

### Expression of recombinant porcine lactoferrin N-lobe in *Pichia methanolica* and its antibacterial activity

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Volume 16 (2007), No. 2, page 288, Figure 2

Instead of:

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1  ATGAAGCTCTTCATCCCCGCCCTGCTGTTCTCGGGACACTTGGACTGTGTCTGGCTGCC 60
1  M K L F I P A L L F L W T L G L C L A 20
61 CCTAAGAAAGGGGTTTCGATGGTGTGTCATATCCACAGCAGAGTATTCAAATGCCGCCAG 120
21 P K K G V R W C V I S T A E Y S K C R Q 40
121 TGGCAATCAAAGATAAGAAGAACTAATCCCATGTTCTGCATAAGGAGGGCTTCTCCCACT 180
41 W Q S K I R R T N P I F C I R R A S P T 60
181 GACTGTATCCGGGCCATCGCGGCAAAAAGGGCAGATGCTGTGACCCTTGATGGTGGTTTG 240
61 D C I R A I A A K R A D A V T L D G G L 80
241 GTGTTTGAAGCAGACCAGTACAAACTGCGGCCGGTAGCAGCGGAGATCTACGGGACAGAA 300
81 V F E A G Q Y K L R P V A A E I Y G T E 100
301 GAGAAATCCCCAAACCTACTATTATGCTGTGGCTGTAGTGAAGAAAGGTTTCAACTTTCAG 360
101 E N P Q T Y Y Y A V A V V K K G F N F Q 120
361 CTGAACCAGCTACAAGGTCGAAAGTCCTGCCACACAGGCCTTGGCAGGTCTGCCGGGTGG 420
121 L N Q L Q G R K S C H T G L G R S A G W 140
421 AATATCCCTATAGGGTTACTTCGCGGGTTC TTGACTGGGCAGGGCCACCTGAGCCCCTC 480
141 N I P I G L L R R F L D W A G P P E P L 160
481 CAGAAAGCTGTGGCCAAATCTTCTCAGAGCTGTGTGCCTCGCCAGATGGAAATGCG 540
161 Q K A V A K F F S Q S C V P C A D G N A 180
541 TATCCCAACCTGTGTGTCAGCTGTGCATAGGGAAAGGAAAGATAAATGTGCTTGTCTCTC 600
181 Y P N L C Q L C I G K G K D K C A C S S 200
601 CAGGAACCGTATTTTGGCTATTCCGGTGCCTTCAACTGTCTGCACAAAGGGATTGGAGAT 660
201 Q E P Y F G Y S G A F N C L H K G I G D 220
661 GTGGCTTTTGTCAAGGAGAGTACAGTGTGAGAACCTGCCACAGAAGGCTGACCGGGAC 720
221 V A F V K E S T V F E N L P Q K A D R D 240
721 AAATACGAGCTACTCTGCCAGACAATACGAAAGCCAGTGGAAGCATTCAGGGAGTGC 780
241 K Y E L L C P D N T R K P V E A F R E C 260
781 CACCTTGCCCGGGTCCCTTCTCATGCTGTGTGGCCCGAAGTGTGAATGGCAAGGAGAAC 840
261 H L A R V P S H A V V A R S V N G K E N 280
841 TCCATCTGGGAGCTTCTTACCAGTCACAGAAAAAGTTTGGAAAAAGCAATCCACAGGAG 900
281 S I W E L L Y Q S Q K K F G K S N P Q E 300
901 TTCCAGCTCTTTGGCTCTCTGGTCAAGCAGAAAGGACCTCTGTTTAGAGATGCTACCATC 960
301 F Q L F G S P G Q K D L L F R D A T I 320
961 GGGTTTTTGAAGATCCCTCAAAGATAGATTCTAAGCTGTACCTGGGCCTCCCGTACCTT 1020
321 G F L K I P S K I D S K L Y L G L P Y L 340
1021 ACTGCCATCCAGGGCCTGA 1038
341 T A I Q G L 346
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Should be corrected version of Figure 2:

ATG AAG CTC TTC ATC CCC GCC CTG CTG TTC CTC GGG ACA CTT GGA CTG TGT CTG GCT GCC	60
<i>M K L F I P A L L F L W T L G L C L A A</i>	20
CCT AAG AAA GGG GTT CGA TGG TGT GTC ATA TCC ACA GCA GAG TAT TCA AAA TGC CGC CAG	120
<i>P K K G V R W C V I S T A E Y S K C R Q</i>	40
TGG CAA TCA AAG ATA AGA AGA ACT AAT CCC ATG <u>TTC</u> TGC ATA AAG AGG OCT TCT CCC ACT	180
<i>W Q S K I R R T N P I F C I R R A S P T</i>	60
GAC TGT ATC CGG GCC ATC GCG GCA AAAAAGG GCA GAT GCT GTG ACC CTT GAT GGT GGT TTG	240
<i>D C I R A I A A K R A D A V T L D G G L</i>	80
GTG TTT GAA CGA GAC CAG TAC AAA CTG GGG CGG GTAGCA GCG GAG ATC TAC GGG ACA GAA	300
<i>V F E A G Q Y K L R P V A A E I Y G T E</i>	100
GAG AAT CCC CAA ACC TAC TAT TAT GCT GTG GCT GTA GTG AAG AAAAGGT TTC AAC TTT CAG	360
<i>E N P Q T Y Y Y A V A V V K K G F N F Q</i>	120
CTG AAC CAG CTA CAAGGT CGA AAG TCC TGC CAC ACA GGC CTT GGC AGG TCT GCC GGG TGG	420
<i>L N Q L Q G R K S C H T G L G R S A G W</i>	140
AAT ATC CCT ATA GGG TTA CTT CCG CGG TTC TTG GAC TGG GCA GGG CCA CCT GAG CCC CTC	480
<i>N I P I G L L R R F L D W A G P P E P L</i>	160
CAG AAA GCT GTG GCC AAA TTC TTC TCT CAG AGC TGT GTG CCC TGC GCA GAT GGAAT GCG	540
<i>Q K A V A K F F S Q S C V P C A D G N A</i>	180
TAT CCC AAC CTG TGT CAG CTG TGC ATA GGG AAA GGG AAA GAT AAA TGT GCT TGT TCC TCC	600
<i>Y P N L C Q L C I G K G K D K C A C S S</i>	200
CAG GAA CCG TAT TTT GGC TAT TCC GGT GCC TTC AAC TGT CTG CAC AAA GGG ATT GGAGAT	660
<i>Q E P Y F G Y S G A F N C L H K G I G D</i>	220
GTG GCT TTT GTC AAG GAG AGT ACA GTG TTT GAG AAC CTG CCACAG AAG OCT GAC CGG GAC	720
<i>V A F V K E S T V F E N L P Q K A D R D</i>	240
AAA TAC GAG CTA CTC TGC CCAGAC AAT ACT CGA AAG CCA GTG GAA GCA TTC AAG GAG TGC	780
<i>K Y E L L C P D N T R K P V E A F R E C</i>	260
CAC CTT GCC CGG GTC CCT TCT CAT GCT GTT GTG GCC CGA AGT GTG AAT GGC AAG GAG AAC	840
<i>H L A R V P S H A V V A R S V N G K E N</i>	280
TCC ATC TGG GAG CTT CTC TAC CAG TCA CAG AAAAAG TTT GGAAAAAGCAAT CCAC AG GAG	900
<i>S I W E L L Y Q S Q K K F G K S N P Q E</i>	300
TTC CAG CTC TTT GGC TCT CCT GGT CAG CAG AAG GAC CTC CTG TTT AGA GAT OCT ACC ATC	960
<i>F Q L F G S P G Q Q K D L L F R D A T I</i>	320
GGG TTT TTG AAG ATC CCC TCA AAG ATA GAT TCT AAG CTG TAC CTG GGC CTC CGG TAC CTT	1020
<i>G F L K I P S K I D S K L Y L G L P Y L</i>	340
ACTGCCATC CAG GGC CTG A	1038
<i>T A I Q G L</i>	346

Figure 2. Sequence of the cloned cDNA encoding the PLfN. The cDNA sequence and the deduced amino acid sequence were shown. The points mutation were underlined and the 17-amino acid signal peptide was written in italics