Appendix A. Supplementary data

Molecular modelling of lactoferrin for food application: Insights from *in silico* techniques

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BLf HLf

₁ _	APRKNVRWCTISQPEWFKCRRWQWRMKKLGAPSITCVRRAFALECIRAIA	50
1	GRŔRSVQŴĊAVŚNPĖATŔĊFQŴQRNMRŔVRGPPVSĊIKŔDSPIQĊĬQĂĬĂ	50
51	EKKADAVTLDGGMVFEAGRDPYKLRPVAAEIYGTKESPQTHYYAVAVVKK	100
51	ENRADAVTLDGGFIYEAGLAPYKLRPVAAEVYGTERQPRTHYYAVAVVKK	100
101	GSNFQLDQLQGRKSCHTGLGRSAGWIIPMGILRPYLSWTESLEPLQGAVA	150
101	GGSFQLNELQGLKSCHTGLRRTAGWNVPIGTLRPFLNWTGPPEPIEAAVA	150
151	KFFSASCVPCIDRQAYPNLCQLCKGEGENQCACSSREPYFGYSGAFKCLQ	200
151	RFFSASCVPGADKGQFPNLCRLCAGTGENKCAFSSQEPYFSYSGAFKCLK	200
201	DGAGDVAFVKETTVFENLPEKADRDQYELLCLNNSRAPVDAFKECHLAQV	250
201	DGAGDVAFIRESTVFEDLSDEAERDEYELLCPDNTRKPVDKFKDCHLARV	250
251	PSHAVVARSVDGKEDLIWKLLSKAQEKFGKNKSRSFQLFGSPPGQRDLLF	300
251	PSHAVVARSVNGKEDAIWNLLRQAQEKFGKDKSPKFQLFGSPSGQKDLLF	300
301	KDSALGFLRIPSKVDSALYLGSRYLTTLKNLRETAEEVKARYTRVVWCAV	350
301	KDSAIGFSRVPPRIDSGLYLGSGYFTAIQNLRKSEEEVAARRARVVWCAV	350
351	GPEEQKKCQQWSQQSGQNVTCATASTTDDCIVLVLKGEADALNLDGGYIY	400
351	GEQELRKCNQWSGLSEGSVTCSSASTTEDCIALVLKGEADAMSLDGGYVY	400
401	TAGKCGLVPVLAENRKSSKHSSLDCVLRPTEGYLAVAVVKKANEGLTW	448
401	TAGKCGLVPVLAENYKSAQSSDPDPNCVDRPVEGYLAVAVVRRSDTSLTW	450
449	NSLKDKKSCHTAVDRTAGWNIPMGLIVNQTGSCAFDEFFSQSCAPGADPK	498
451	NSVKGKKSCHTAVDRTAGWNIPMGLLFNQTGSCKFDEYFSQSCAPGSDPA	500
499	SRLCALCAGDDQGLDKCVPNSKEKYYGYTGAFRCLAEDVGDVAFVKNDTV	548
501	SNLCALCIGDEEGENKCVPNSNERYYGYTGAFRCLAENAGDVAFVKDVTV	550
549	WENTNGESTADWAKNLNREDFRLLCLDGTRKPVTEAQSCHLAVAPNHAVV	598
551	LQNTDGNNNEAWAKDLKLADFALLCLDGKRKPVTEARSCHLAMAPNHAVV	600
599	SRSDRAAHVKQVLLHQQALFGKNGKNCPDKFCLFKSETKNLLFNDNTECL	648
601	SRMDKVERLKQVLLHQQAKFGRNGSDCPDKFCLFQSETKNLLFNDNTECL	650
649	AKLGGRPTYEEYLGTEYVTAIANLKKCSTSPLLEACAFLTR 689	
651	ARLHGKTTYEKYLGPQYVAGITNLKKCSTSPLLEACEFLRK 691	

Figure S1. Comparison of the amino acid sequencing results for bovine lactoferrin (BLf) and human lactoferrin (HLf). Symbols adjoining the single-letter amino acid codes have following meanings: '|' = identical residue; ':' = BLOSUM matrix score > 0; and '.' = BLOSUM matrix score ≤ 0.



Figure S2. A cartoon representation of the stretching of bonds



Figure S3. A cartoon representation of the bond angle calculation



Figure S4. A cartoon representation of the dihedral angle



Figure S5. A cartoon representation of the improper dihedral angle



Figure S6. A cartoon representation of Lennard-Jones interactions



Figure S7. A cartoon representation of electrostatic interactions



Figure S8. A cartoon representation of the Urey-Bradley potential



Figure S9. A cartoon representation of hydrogen bonds



Figure S10. A 3D illustration of a periodic boundary condition