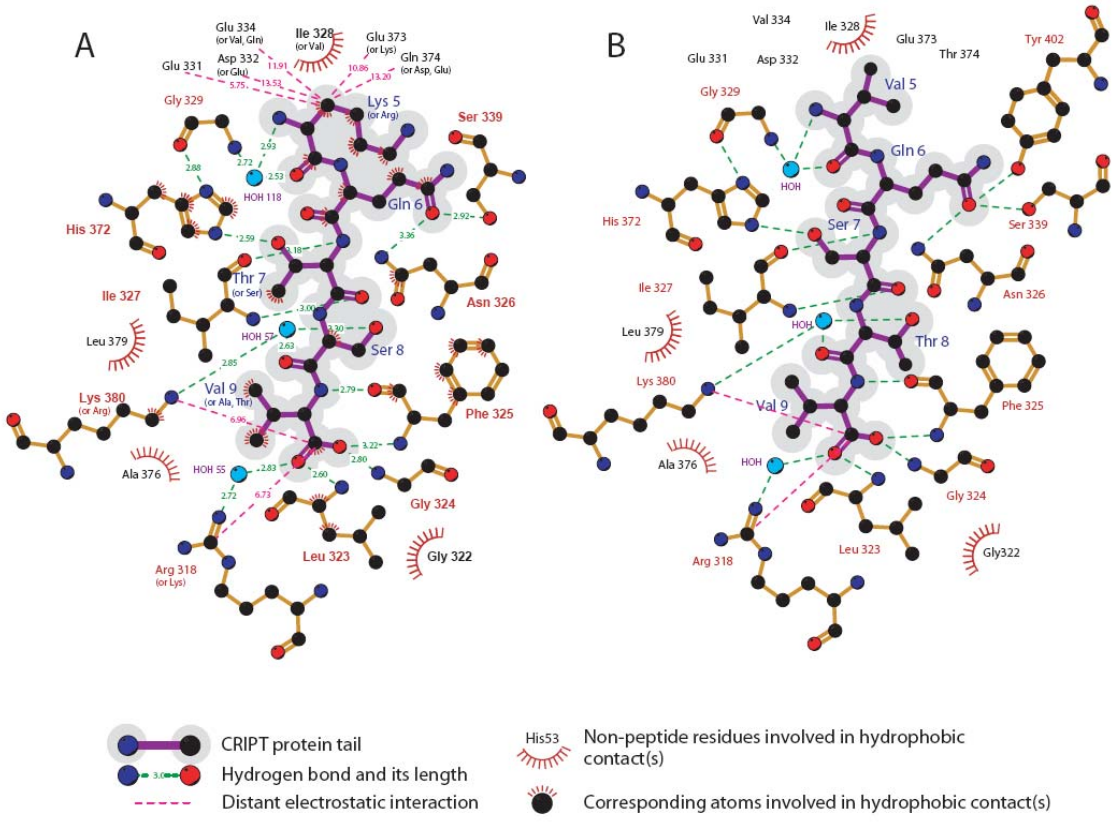
A
Glu 334 (or Val, Gln)
Ile 328 (or Val)
Glu 373 (or Lys)
Glu 331
Asp 332 (or Glu)
Gln 374 (or Asp, Glu)
Gly 329
Lys 5 (or Arg)
Ser 339
HOH 118
Gln 6
His 372
Thr 7 (or Ser)
Ile 327
Asn 326
Leu 379
HOH 57
Ser 8
Val 9 (or Ala, Thr)
Lys 380 (or Arg)
Phe 325
Ala 376
HOH 55
Gly 324
Leu 323
Gly 322
Arg 318 (or Lys)
B
Val 334
Ile 328
Glu 373
Tyr 402
Glu 331
Asp 332
Thr 374
Gly 329
Val 5
HOH
Gln 6
His 372
Ser 7
Ser 339
Asn 326
Ile 327
Leu 379
HOH
Thr 8
Lys 380
Val 9
Phe 325
Ala 376
HOH
Gly 324
Arg 318
Leu 323
Gly322
CRIPT protein tail
Hydrogen bond and its length
Distant electrostatic interaction
His53
Non-peptide residues involved in hydrophobic contact(s)
Corresponding atoms involved in hydrophobic contact(s)