

## hbond.lol

270 TYR H	280 TYR O	1.8
270 TYR N	280 TYR O	2.7
271 HIS H	340 GLY O	1.8
271 HIS N	340 GLY O	2.7
272 VAL H	278 LEU O	1.8
272 VAL N	278 LEU O	2.7
273 LEU H	331 SER OG	1.8
273 LEU N	331 SER OG	2.7
274 VAL H	276 VAL O	1.8
274 VAL N	276 VAL O	2.7
277 LYS H	296 ARG O	1.8
277 LYS N	296 ARG O	2.7
279 VAL H	276 HIS O	1.8
279 VAL N	276 HIS O	2.7
300 ASP H	297 SER OG	1.8
300 ASP N	297 SER OG	2.7
301 ALA H	297 SER O	1.8
301 ALA N	297 SER O	2.7
302 ILE H	298 ARG O	1.8
302 ILE N	298 ARG O	2.7
303 ASN H	299 ALA O	1.8
303 ASN N	299 ALA O	2.7
304 LEU H	300 ASP O	1.8
304 LEU N	300 ASP O	2.7
305 ALA H	301 ALA O	1.8
305 ALA N	301 ALA O	2.7
306 GLN H	302 ILE O	1.8
306 GLN N	302 ILE O	2.7
307 ALA H	303 ASN O	1.8
307 ALA N	303 ASN O	2.7
308 ILE H	304 LEU O	1.8
308 ILE N	304 LEU O	2.7
309 LEU H	305 ALA O	1.8
309 LEU N	305 ALA O	2.7
310 ALA H	306 GLN O	1.8
310 ALA N	306 GLN O	2.7
322 GLU H	319 SER OG	1.8
322 GLU H	319 SER OG	2.7
323 PHE H	319 SER O	1.8
323 PHE H	319 SER O	2.7
324 VAL H	320 LEU O	1.8
324 VAL N	320 LEU O	2.7
325 GLN H	321 ASP O	1.8
325 GLN N	321 ASP O	2.7
326 VAL H	323 PHE O	1.8
326 VAL N	323 PHE O	2.7
327 VAL H	323 PHE O	1.8
327 VAL N	323 PHE O	2.7
328 ARG H	324 VAL O	1.8
328 ARG N	324 VAL O	2.7
330 PHE H	326 VAL O	1.8
330 PHE N	326 VAL O	2.7
331 SER H	327 VAL O	1.8
331 SER N	327 VAL O	2.7
332 GLU H	273 LEU O	1.8
332 GLU N	273 LEU O	2.7
333 CYS H	331 SER OG	1.8
333 CYS N	331 SER OG	2.7
341 ASP H	338 ARG O	1.8
341 ASP N	338 ARG O	2.7
342 LEU H	269 PHE O	1.8
342 LEU N	269 PHE O	2.7
358 ALA H	354 PHE O	1.8

## hbond.upl

270 TYR H	280 TYR O	2.6
270 TYR N	280 TYR O	3.5
271 HIS H	340 GLY O	2.6
271 HIS N	340 GLY O	3.5
272 VAL H	278 LEU O	2.6
272 VAL N	278 LEU O	3.5
273 LEU H	331 SER OG	2.6
273 LEU N	331 SER OG	3.5
274 VAL H	276 VAL O	2.6
274 VAL N	276 VAL O	3.5
277 LYS H	296 ARG O	2.6
277 LYS N	296 ARG O	3.5
279 VAL H	276 HIS O	2.6
279 VAL N	276 HIS O	3.5
300 ASP H	297 SER OG	2.6
300 ASP N	297 SER OG	3.5
301 ALA H	297 SER O	2.6
301 ALA N	297 SER O	3.5
302 ILE H	298 ARG O	2.6
302 ILE N	298 ARG O	3.5
303 ASN H	299 ALA O	2.6
303 ASN N	299 ALA O	3.5
304 LEU H	300 ASP O	2.6
304 LEU N	300 ASP O	3.5
305 ALA H	301 ALA O	2.6
305 ALA N	301 ALA O	3.5
306 GLN H	302 ILE O	2.6
306 GLN N	302 ILE O	3.5
307 ALA H	303 ASN O	2.6
307 ALA N	303 ASN O	3.5
308 ILE H	304 LEU O	2.6
308 ILE N	304 LEU O	3.5
309 LEU H	305 ALA O	2.6
309 LEU N	305 ALA O	3.5
310 ALA H	306 GLN O	2.6
310 ALA N	306 GLN O	3.5
322 GLU H	319 SER OG	2.6
322 GLU H	319 SER OG	3.5
323 PHE H	319 SER O	2.6
323 PHE H	319 SER O	3.5
324 VAL H	320 LEU O	2.6
324 VAL N	320 LEU O	3.5
325 GLN H	321 ASP O	2.6
325 GLN N	321 ASP O	3.5
326 VAL H	323 PHE O	2.6
326 VAL N	323 PHE O	3.5
327 VAL H	323 PHE O	2.6
327 VAL N	323 PHE O	3.5
328 ARG H	324 VAL O	2.6
328 ARG N	324 VAL O	3.5
330 PHE H	326 VAL O	2.6
330 PHE N	326 VAL O	3.5
331 SER H	327 VAL O	2.6
331 SER N	327 VAL O	3.5
332 GLU H	273 LEU O	2.6
332 GLU N	273 LEU O	3.5
333 CYS H	331 SER OG	2.6
333 CYS N	331 SER OG	3.5
341 ASP H	338 ARG O	2.6
341 ASP N	338 ARG O	3.5
342 LEU H	269 PHE O	2.6
342 LEU N	269 PHE O	3.5
358 ALA H	354 PHE O	2.6

358 ALA N	354 PHE O	2.7	358 ALA N	354 PHE O	3.5
361 LEU H	358 ALA O	1.8	361 LEU H	357 VAL O	2.6
361 LEU N	358 ALA O	2.7	361 LEU N	357 VAL O	3.5
364 GLY H	281 ARG O	1.8	364 GLY H	281 ARG O	2.6
364 GLY N	281 ARG O	2.7	364 GLY N	281 ARG O	3.5
365 GLU H	362 LYS O	1.8	365 GLU H	362 LYS O	2.6
365 GLU N	362 LYS O	2.7	365 GLU N	362 LYS O	3.5
367 SER H	279 ILE O	1.8	367 SER H	279 ILE O	2.6
367 SER N	279 ILE O	2.7	367 SER N	279 ILE O	3.5
370 VAL H	277 HIS O	1.8	370 VAL H	277 HIS O	2.6
370 VAL N	277 HIS O	2.7	370 VAL N	277 HIS O	3.5
372 THR H	275 GLY O	1.8	372 THR H	275 GLY O	2.6
372 THR N	275 GLY O	2.7	372 THR N	275 GLY O	3.5
375 GLY H	272 THR O	1.8	375 GLY H	272 THR O	2.6
375 GLY N	272 THR O	2.7	375 GLY N	272 THR O	3.5
376 VAL H	274 VAL O	1.8	376 VAL H	274 VAL O	2.6
376 VAL N	274 VAL O	2.7	376 VAL N	274 VAL O	3.5
377 HIS H	270 VAL O	1.8	377 HIS H	270 VAL O	2.6
377 HIS N	270 VAL O	2.7	377 HIS N	270 VAL O	3.5
378 LEU H	272 VAL O	1.8	378 LEU H	272 VAL O	2.6
378 LEU N	272 VAL O	2.7	378 LEU N	272 VAL O	3.5
379 ILE H	267 SER OG	1.8	379 ILE H	267 SER OG	2.6
379 ILE N	267 SER OG	2.7	379 ILE N	267 SER OG	3.5
380 TYR H	270 TYR O	1.8	380 TYR H	270 TYR O	2.6
380 TYR N	270 TYR O	2.7	380 TYR N	270 TYR O	3.5
381 ARG H	265 GLU O	1.8	381 ARG H	265 GLU O	2.6
381 ARG N	265 GLU O	2.7	381 ARG N	265 GLU O	3.5
382 VAL H	268 HIS O	1.8	382 VAL H	268 HIS O	2.6
382 VAL N	268 HIS O	2.7	382 VAL N	268 HIS O	3.5
			286 ALA QB	336 ALA QB	3.8