

In silico study of 4-amino substituted-7-chloroquinoline derivatives as *Plasmodium falciparum* lactate dehydrogenase inhibitors

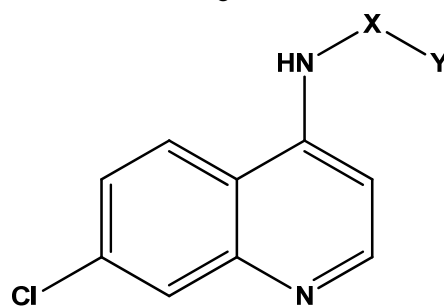
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Received 28 April 2023; revised 25 April 2024

Supplementary Data

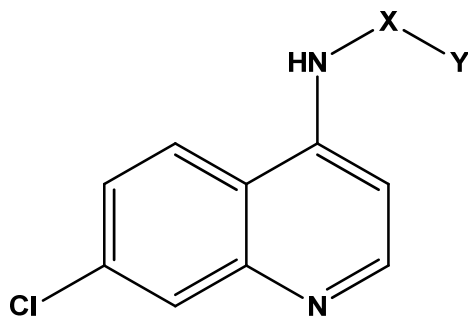
Table 1 — Amino acids residues interacting with 4-amino substituted-7-chloroquinoline



Compounds 1-25

Comp. No	X	Y	Amino acids residues interacting with 4-amino substituted-7-chloroquinoline
1	-CH ₂ -CO-	Phenyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Ile119 Glu122 Ile123
2	-(CH ₂) ₂ -CO-	Phenyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Pro114 Ile119 Glu122 Ile123
3	-(CH ₂) ₃ -CO-	Phenyl	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Lys56 Tyr85 Ala98 Gly99 Ile119 Glu122 Ile123
4	-(CH ₂) ₄ -CO-	Phenyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Ile119 Glu122 Ile123 His126
5	-(CH ₂) ₅ -CO-	Phenyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Lys 118 Ile119 Glu122 Ile123
6	-(CH ₂) ₆ -CO-	Phenyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Pro114 Leu115 Asn117 Lys118 Ile119 Glu122 Ile123
7	-CH ₂ -CO-	2-Furyl	Gly27 Ser28 Gly29 Met30 Ile31 Gly32 Asp53 Thr97 Ala98 Gly99 Thr101 Val138 Thr139 Asn140 Ser245 Pro246 Tyr247
8	-(CH ₂) ₂ -CO-	2-Furyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Lys118 Ile119 Glu122 Ile123
9	-(CH ₂) ₃ -CO-	2-Furyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Lys118 Ile119 Glu122 Ile123
10	-(CH ₂) ₄ -CO-	2-Furyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Lys118 Ile119 Glu122 Ile123
11	-(CH ₂) ₅ -CO-	2-Furyl	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Lys118 Ile119 Glu122 Ile123
12	-(CH ₂) ₆ -CO-	2-Furyl	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Pro114 Leu115 Asn117 Lys118 Ile119 Glu122 Ile123

(Contd.)

Table 1 — Amino acids residues interacting with 4-amino substituted-7-chloroquinoline (*Contd.*)

Compounds 1-25

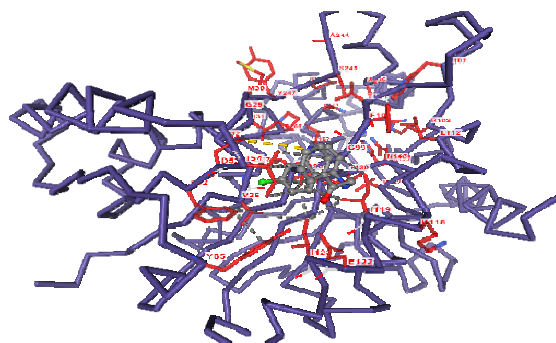
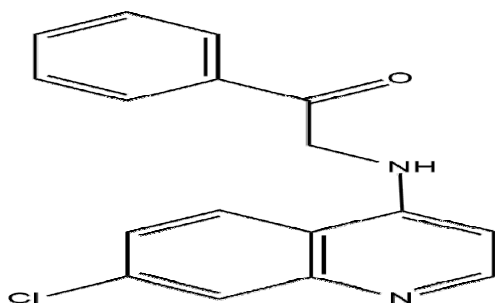
Comp. No	X	Y	Amino acids residues interacting with 4-amino substituted-7-chloroquinoline
13	-CH ₂ -CO-	2-Thiophene	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Thr101 Pro114 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
14	-(CH ₂) ₂ -CO-	2-Thiophene	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
15	-(CH ₂) ₃ -CO-	2-Thiophene	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
16	-(CH ₂) ₄ -CO-	2-Thiophene	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
17	-(CH ₂) ₅ -CO-	2-Thiophene	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
18	-(CH ₂) ₆ -CO-	2-Thiophene	Val26 Gly27 Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Leu115Asn117 Lys118 Ile119 Glu122 Ile123
19	-CH ₂ -CO-	2-Azole	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Leu115Lys118 Ile119 Glu122 Ile123
20	-(CH ₂) ₂ -CO-	2-Azole	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Ile119 Glu122Ile123
21	-(CH ₂) ₃ -CO-	2-Azole	Val26 Gly27 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Leu115Lys118 Ile119 Glu122 Ile123
22	-(CH ₂) ₄ -CO-	2-Azole	Val26 Gly27Ser28 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Leu115Lys118 Ile119 Glu122 Ile123
23	-(CH ₂) ₅ -CO-	2-Azole	Val26 Gly27 Ser28Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Leu115Lys118 Ile119 Glu122 Ile123
24	-(CH ₂) ₆ -CO-	2-Azole	Val26 Gly27 Ser28Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Leu115Lys118 Ile119 Glu122 Ile123
25	-(CH)-CH ₃ -	3-NH ₂ Phenyl	Val26 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Ile119 Glu122Ile123
26	Chloroquine		Val26 Phe52 Asp53 Ile54 Tyr85 Ala98 Gly99 Phe100 Ile119 Glu122Ile123

Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives

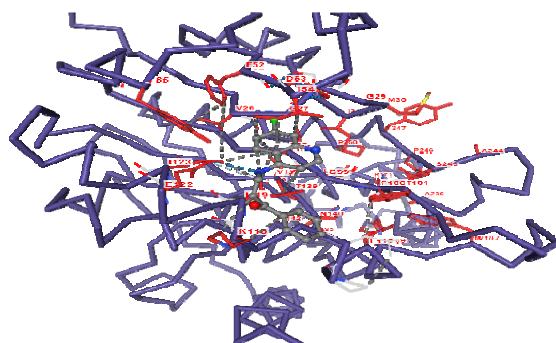
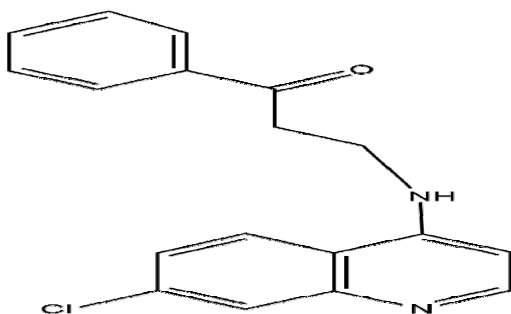
Sr. No. Structure

Docking images

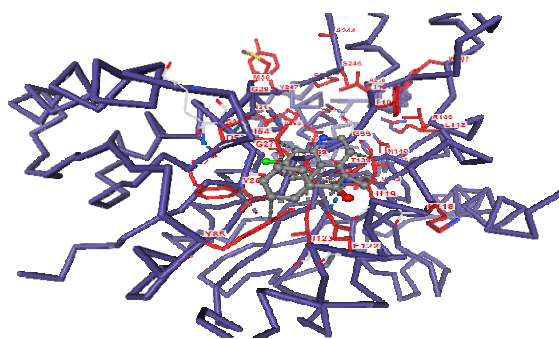
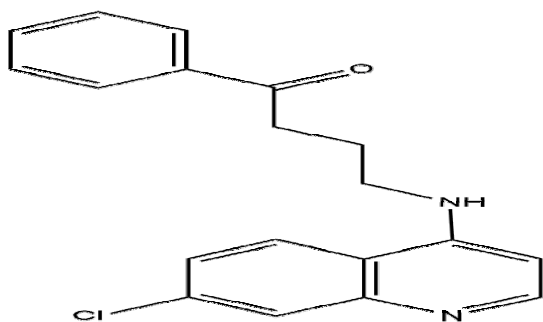
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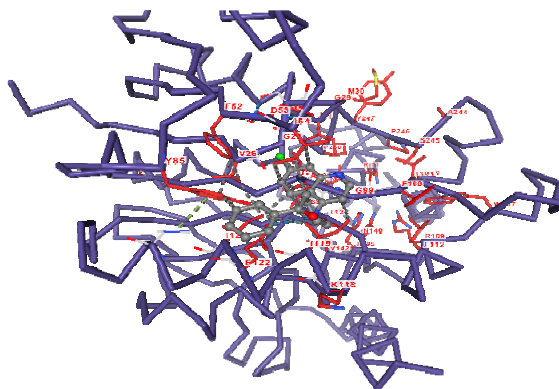
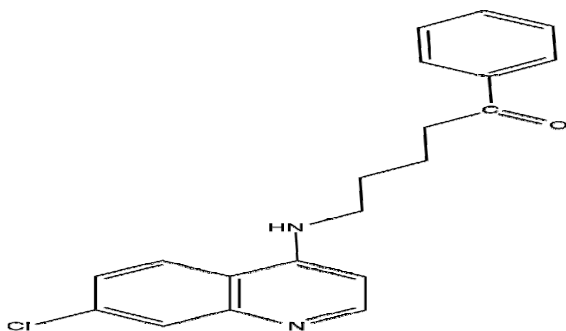
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4



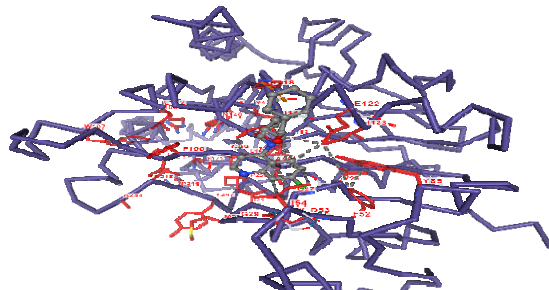
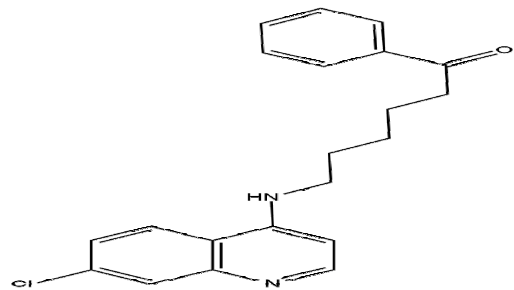
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Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives (*Contd.*)

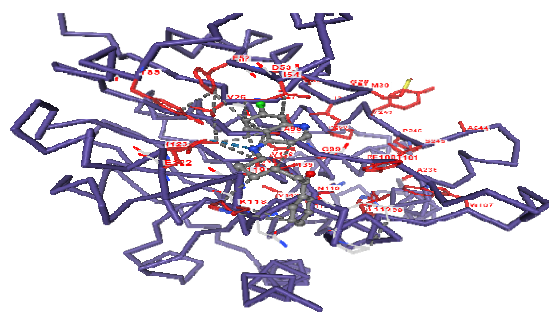
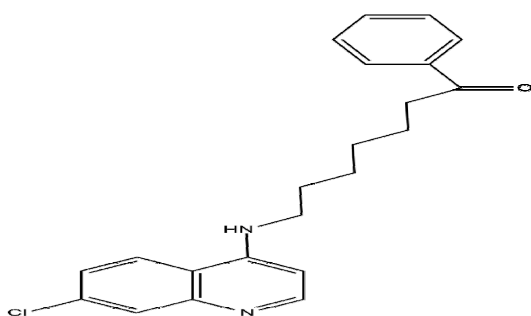
Sr. No. Structure

Docking images

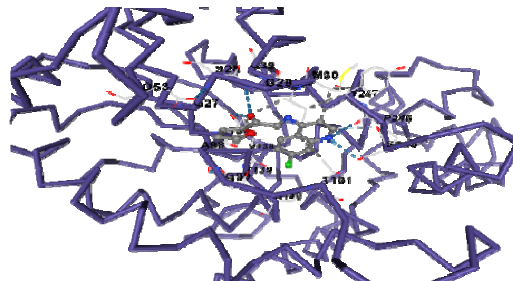
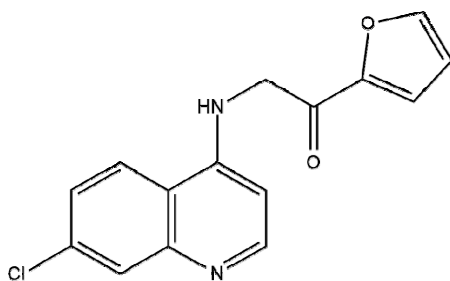
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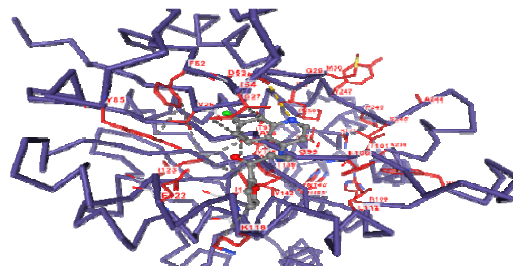
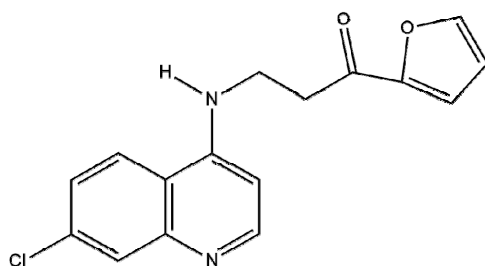
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7



8



9

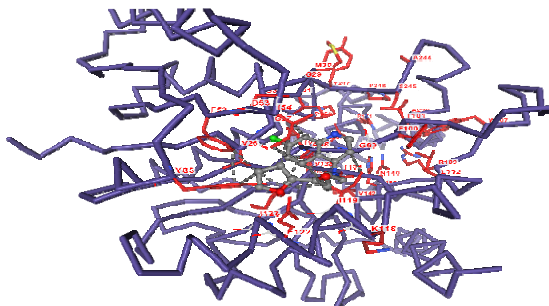
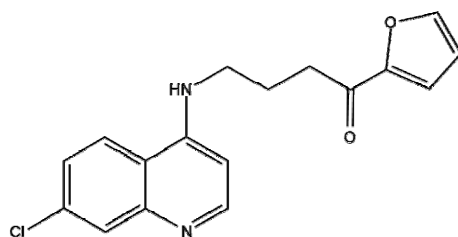
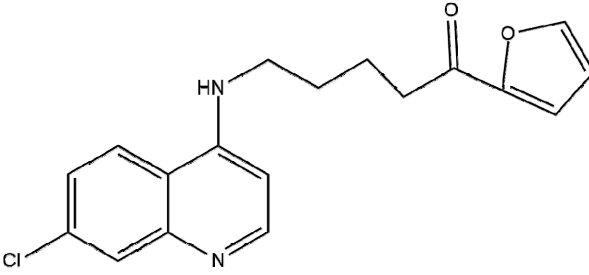
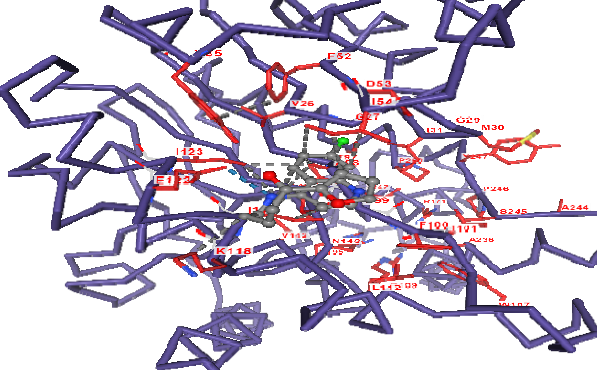
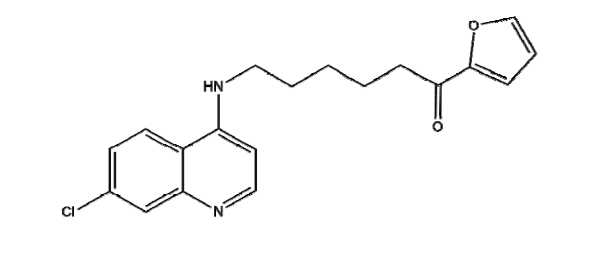
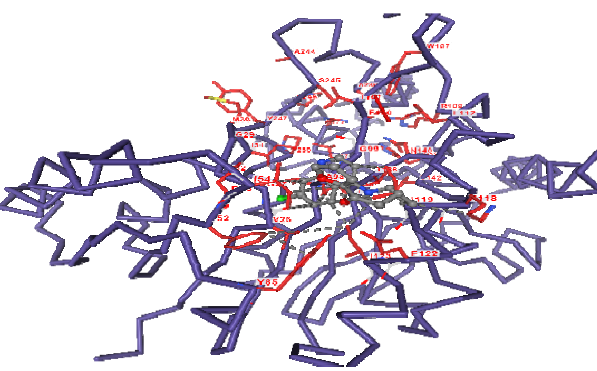
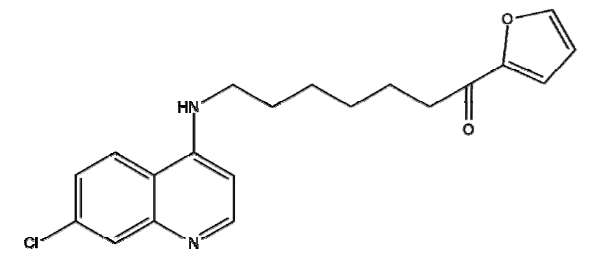
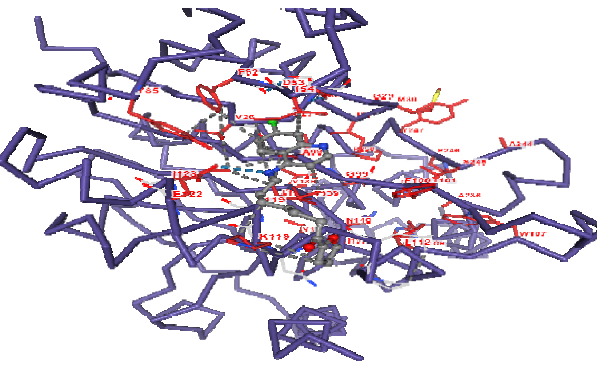
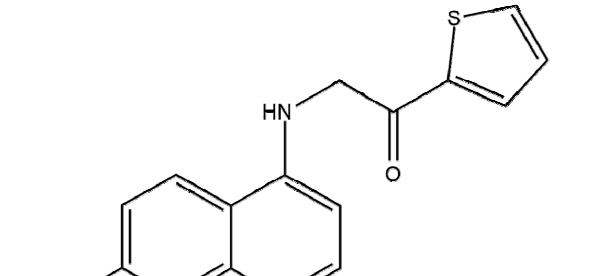
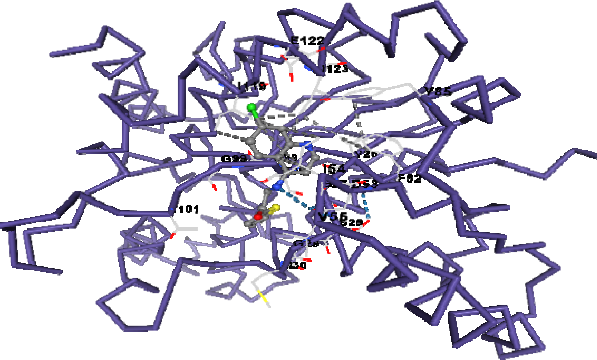
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Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives (*Contd.*)

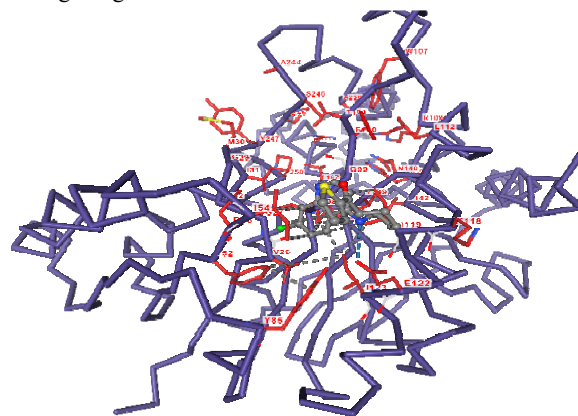
Sr. No.	Structure	Docking images
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11		
12		
13		

(*Contd.*)

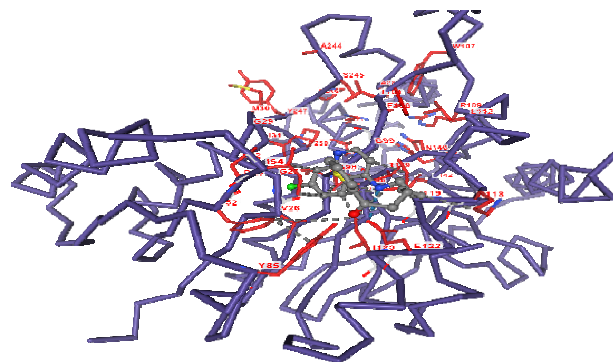
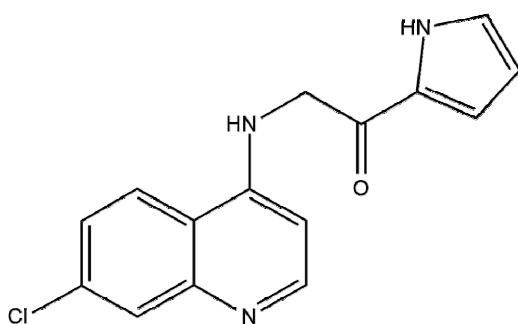
Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives (*Contd.*)

Sr. No. Structure
18

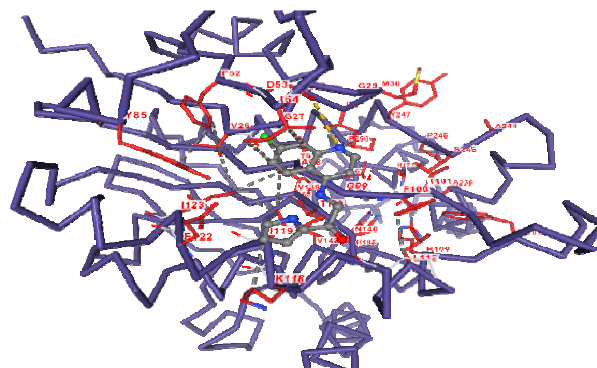
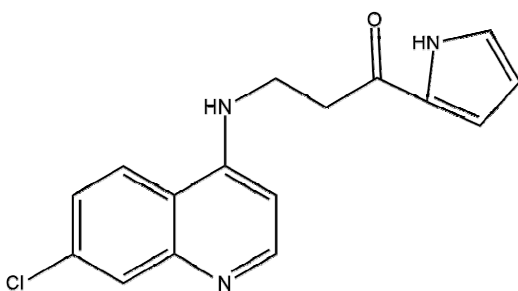
Docking images



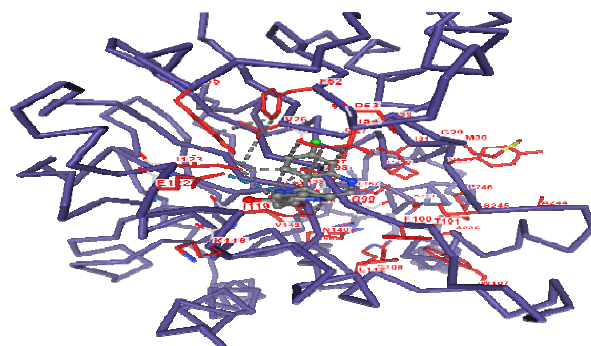
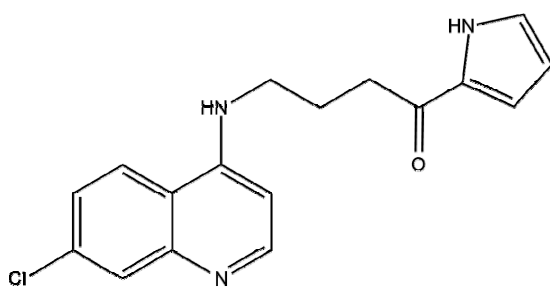
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20



21



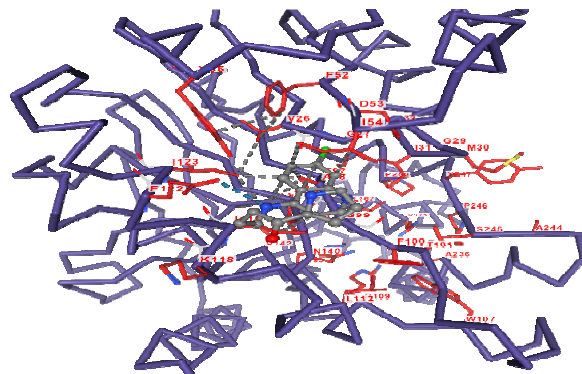
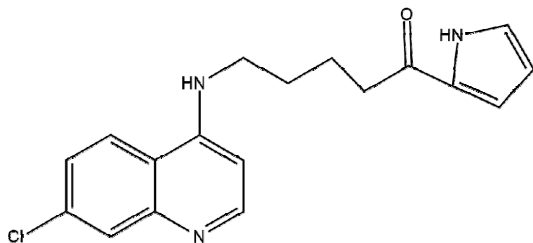
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Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives (*Contd.*)

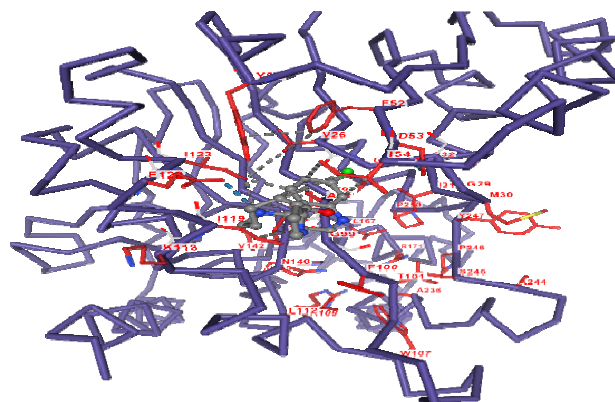
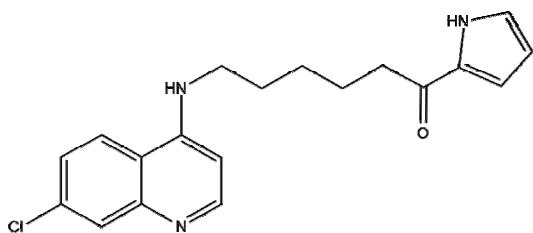
Sr. No. Structure

Docking images

22



23



24

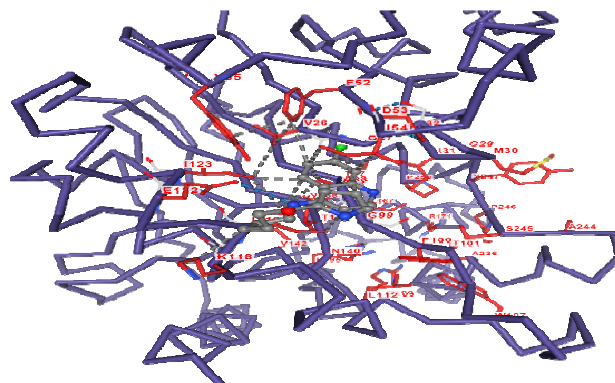
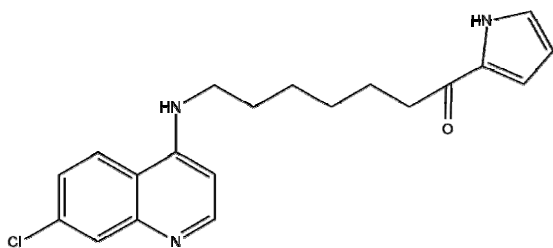
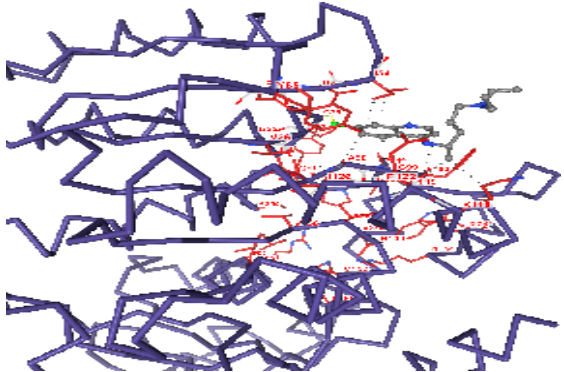


Table 2 — Docking images of designed 4-aminosubstituted-7-chloroquinoline derivatives (*Contd.*)

Sr. No.	Structure	Docking images
26	Chloroquine	 A 3D molecular docking image showing the interaction of Chloroquine (represented by a blue stick model) with the active site of the Plasmodium falciparum Lactate Dehydrogenase (LDH) enzyme. The protein backbone is shown in a light grey wireframe. Red lines and labels indicate specific interactions between the ligand and the enzyme's residues, such as hydrogen bonds and hydrophobic contacts. The ligand is positioned within the enzyme's binding pocket, with its quinoline ring system interacting with several residues.