

## **BAB V**

### **KESIMPULAN DAN SARAN**

#### **A. Kesimpulan**

Pertama, nilai afinitas dari 26 senyawa turunan 3-aminopiridin-2(1H)-on tidak berkorelasi dengan aktivitas *in vitro*.

Kedua, pola interaksi 26 senyawa turunan 3-aminopiridin-2(1H)-on terhadap enzim RT HIV-1 dengan melibatkan asam amino yang berbeda-beda berada pada empat daerah yang berbeda, yaitu:

- a. Daerah penghambatan pertama menempati daerah yang sama seperti daerah penghambatan ligan asli dan ditempati oleh 17 senyawa turunan 3-aminopiridin-2(1H)-on.
- b. Daerah penghambatan kedua ditempati oleh tiga senyawa turunan 3-aminopiridin-2(1H)-on dengan jarak sekitar 12,51 $\text{\AA}$  dengan daerah penghambatan pertama.
- c. Daerah penghambatan ketiga ditempati oleh tiga senyawa turunan 3-aminopiridin-2(1H)-on dengan jarak sekitar 10,34 $\text{\AA}$  dengan daerah penghambatan pertama
- d. Daerah penghambatan keempat ditempati oleh tiga senyawa turunan 3-aminopiridin-2(1H)-on dengan jarak sekitar 49,14 $\text{\AA}$  dari daerah penghambatan.

## B. Saran

Pertama, perlu dilakukan penelitian *docking* molekuler senyawa turunan 3-aminopiridin-2(1H)-on terhadap enzim RT HIV-1 dengan *binding site* dan metode yang berbeda.

Kedua, pemilihan pose dilakukan lebih mendetail pada setiap posenya dengan memperhatikan reproduksibilitas yang paling baik.

Ketiga, menggunakan sistem operasi linux.

.

## DAFTAR PUSTAKA

- Alcaro S, Artese A, Ceccherini-Silberstein F, Chiarella V, Dimonte S, Ortuso F, Perno CF. 2010. Computational analysis of human immunodeficiency virus (HIV) type-1 reverse transcriptase crystallographic models based on significant conserved residues found in highly active antiretroviral therapy (HAART)-treated patients. *Current Medicinal Chemistry* 17:290-308.
- AutoDock Vina. <http://vina.scripps.edu/>.
- Balzarini J. 2004. Current status of the non-nucleoside reverse transcriptase inhibitors of human immunodeficiency virus type 1. *Current Topics in Medicinal Chemistry* 4:921-944.
- Baratawidjaja KG, Rengganis I. 2010. *Imunologi Dasar Edisi ke-9*. Jakarta: Balai Penerbit FKUI.
- Camarasa MJ, Velàzquez S, San-Fèlix A, Pèrez-Pèrez MJ, Gago F. 2006. Dimerization inhibitors of HIV-1 reverse transcriptase, PR and integrase: a single mode of inhibition for the three HIV enzymes? *Antiviral Research* 71:260-267.
- Cramer CJ. 2004. *Essentials of Computational Chemistry Second Edition*. England: John Wiley & Sons.
- De Clercq E. 2002. Strategies in the design of antiviral drugs. *Nat. Rev. Drug Discovery* 1:13–25.
- De Clercq E. 2009. Anti-HIV drugs: 25 compounds approved within 25 years after the discovery of HIV. *International Journal of Antimicrobial Agents* 33:307-320.
- Esnouf RM, Ren J, Hopkins AL, Ross CK, Jones EY, Stammers DK, Stuart DI. 1997. Unique features in the structure of the complex between HIV-1 reverse transcriptase and the bis(heteroaryl)piperazine (BHAP) U-90152 explain resistance mutations for this nonnucleoside inhibitor. *Proc. Natl. Acad. Sci.* 94:3984-3989.
- Gervay-Hague J. 2006. Current developments in HIV chemotherapy. *ChemBioChem* 1:16–29.
- Gotte M, Li X, Wainberg MA. 1999. HIV-1 reverse transcription: a brief overview focused on structure-function relationships among molecules

- involved in initiation of the reaction. *Arch. Biochem. Biophys.* 365:199-210.
- Himmel DM, Das K, Clark AD, Jr, Hughes SH, Benjahad A, Oumouch S, Guillemont J, Coupa S, Poncelet A, Csoka I, Meyer C, Andries K, Nguyen CH, Grierson DS, Arnold E. 2005. Crystal structures for HIV-1 reverse transcriptase in complexes with three piridinon derivatives: a new class of non-nucleoside reverse transcriptase inhibitor effective against a broad range of drug-resistant strains. *J. Med. Chem.* 48:7582-7591.
- Himmel DM, Das K, Clark AD, Jr, Hughes SH, Benjahad A, Oumouch S, Guillemont J, Coupa S, Poncelet A, Csoka I, Meyer C, Andries K, Nguyen CH, Grierson DS, Arnold E. 2005. Crystal structures for HIV-1 reverse transcriptase in complexes with three piridinon derivatives: a new class of non-nucleoside reverse transcriptase inhibitor effective against a broad range of drug-resistant strains [2BE2]. *J. Med. Chem.* 48:7582-7591. <http://pdb.org>. [7 Oktober 2013].
- Hoffman JM, Wai JS, Thomas CM, Levin RB, O'Brien JA, Goldman ME. 1992. Synthesis and evaluation of 2-pyridinone derivatives as HIV-1 specific reverse transcriptase inhibitors. 1. Phthalimidoalkyl and -alkylamino analogues. *J. Med. Chem.* 35:3784-3791.
- Hoffman JM, Smith AM, Rooney CS, Fisher TE, Wai JS, Thomas CM, Bamberger DL, Barnes JL, Williams TM, Jones JH, Olson BD, O'Brien JA, Goldman ME, Nunberg JH, Quintero JC, Schleif WA, Emini EA, Anderson PS. 1993. Synthesis and evaluation of 2-piridinon derivatives as HIV-1-specific reverse transcriptase inhibitors. 4. 3-[2-(Benzoxazol-2-yl)ethyl]-5-ethyl-6-methylpyridin-2(1H)-on and analogues. *J. Med. Chem.* 36:953-966.
- Julias JG, McWilliams MJ, Sarafianos SG, Alvord WG, Arnold E, Hughes SH. 2003. Mutation of amino acids in the connection domain of human immunodeficiency virus type 1 reverse transcriptase that contact the template-primer affects RNase H activity. *J. Virol.* 15:8548-8554.
- Kohlstaedt LA, Wang J, Friedman JM, Rice PA, Steitz TA. 1992. Crystal structure at 3.5 Å resolution of HIV-1 reverse transcriptase complexed with an inhibitor. *Science* 256:1783-1790.
- Kumar V, Cotran RS, Robbins SL. 2007. *Buku Ajar Patologi*. Ed ke-7. Volume ke-1. Prasetyo A, Pendit BU, Priliono T, penerjemah; Asroruddin M, Hartanto H, Darmaniah N, editor. Jakarta: EGC. Terjemahan dari: *Robbins Basic Pathology 7<sup>th</sup> ed.*

- Medina-Franco JL, Morales SR, Gordiano CJ, Campos AH, Castillo R. 2004. Docking-based CoMFA and CoMSIA studies of non-nucleoside reverse transcriptase inhibitors of the pyridinone derivative type. *Journal of Computer-Aided Molecular Design* 18:345-360.
- Mehellou Y, Clerq E de. 2009. Twenty-six years of anti-HIV drug discovery: where do we stand and where do we go?. *J. Med. Chem* XXX(XX).
- Miyasaka C, Walker RT, Tanaka H, Stammers DK, Stuart DI. 1996. Complexes of HIV-1 reverse transcriptase with inhibitors of the HEPT series reveal conformational changes relevant to the design of potent non-nucleoside inhibitors. *J. Med. Chem.* 39:1589-1600.
- Morris GM, Lim-Wilby M. 2008. Molecular Docking. Di dalam: Kukol A, editor. *Molecular Modeling of Proteins*. Totowa: Humana Press. hlm 365-382.
- Mulky A, Sarafianos SG, Jia Y, Arnold E, Kappes JC. 2005. Identification of amino acid residues in the human immunodeficiency virus type-1 reverse transcriptase tryptophan-repeat motif that are required for subunit interaction using infectious virions. Intrinsic tryptophan fluorescence of bovine liver adenosine kinase, characterization of ligand binding sites and conformational changes. *J. Mol. Biol.* 349:673-684.
- Playfair JHL, Chain PM. 2012. *At a Glance Imunologi edisi kesembilan*. Winardini, penerjemah; Astikawati R, editor. Jakarta: Erlangga. Terjemahan dari: *Immunology At a Glance Ninth Edition*.
- Putz MV, Lazea M, Putz AM, Seiman CD. 2011. Introducing catastrophe-QSAR. Application on modeling molecular mechanism of pyridinone derivative-type HIV non-nucleoside reverse transcriptase inhibitors. *Int. J. Mol. Sci.* 12:9533-9569.
- Ramachandran KI, Deepa G, Namboori K. 2008. *Computational Chemistry and Molecular Modeling*. Berlin: Springer.
- Ravichandran S, Veerasamy R, Raman S, Krishnan PN, Agrawal RK. 2008. An overview on HIV-1 reverse transcriptase inhibitors. *Digest Jurnal of Nanomaterials and Biostuctures* 3:171-187.
- Saari WS, Wai JS, Fisher TE, Thomas CM, Hoffman JM, Rooney CS, Smith AM, Jones JH, Bamberger DL, Goldman ME, O'Brien JA, Nunberg JH, Quintero JC, Schleif WA, Emini EA, Anderson PS. 1992. Synthesis and evaluation of 2-pyridinon derivatives as HIV-1-specific reverse transcriptase inhibitors: 2. analogues of 3-amioypyridin-2(1H)-on. *J. Med. Chem* 35:3792-3802.

- Sarafianos SG, Marchand B, Das K, Himmel D, Parniak MA, Hughes SH, Arnold E. 2009. Structure and function of HIV-1 reverse transcriptase: molecular mechanisms of polymerization and inhibition. *J Mol Biol* 385(3):693-713.
- Schaal W. 2002. Computational studies of HIV-1 PR inhibitor [Disertasi]. Sweden: Department of Medicinal Chemistry, Uppsala University.
- Seeliger D, Groot BL de. 2010. Ligand docking and binding site analysis with PyMol and Autodock/Vina. *J Comput Aided Mol* 24:417-422.
- Siswandono, Soekardjo B., editor. 1995. *Kimia Medisinal*. Ed. ke-1. Surabaya: Universitas Airlangga. hlm 190.
- Tachedjian G, Aronson HEG, Goff SP. 2000. Analysis of mutations and suppressors affecting interactions between the subunits of the HIV type 1 reverse transcriptase. *Proc. Natl. Acad. Sci. U.S.A.* 97:6334-6339.
- Tome C. 2013. Human immunodeficiency virus structure. <http://mappingignorance.org> [21 Oktober 2013].
- Trott O, Olson AJ. 2010. AutoDock Vina: improving the speed and accuracy of docking with a new scoring function, efficient optimization, and multithreading. *J Comp Chem* 31:455–461.
- Unadi YC. 2008. perancangan pentapeptida siklis sebagai inhibitor neuramindase virus H5N1 melalui docking dan simulasi dinamika molekul [Tesis]. Jakarta: Fakultas Matematika dan Ilmu Pengetahuan Alam, Universitas Indonesia.
- Wallace AC, Laskowski RA, Thornton JM. 1995. LIGPLOT: a program to generate schematic diagrams of protein-ligand interaction. *Protein Eng* 8(2):127-34.
- Wolff ME. *Asas-asas Kimia Medisinal*. Edisi Keempat. Muljadi, Sabikis, Sumarno, penerjemah; Sasmito, editor. Yogyakarta: Gadjah Mada University Press. Terjemahan dari: *The Basis of Medicinal Chemistry*.
- Yeturu K, Chandra N. 2008. PocketMatch: a New Algorithm to Compare Binding Sites in Protein Structures. *BMC Bioinformatics* 9:543.
- Yuwono T. 2007. *Biologi Molekular*. Jakarta: Erlangga. Hlm 31.

L

A

M

P

I

R

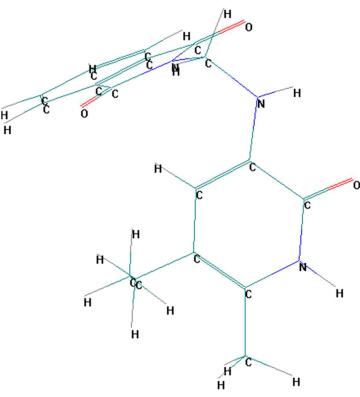
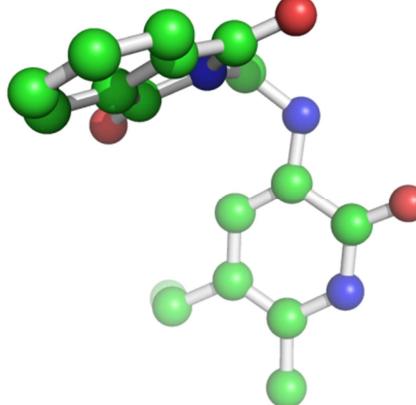
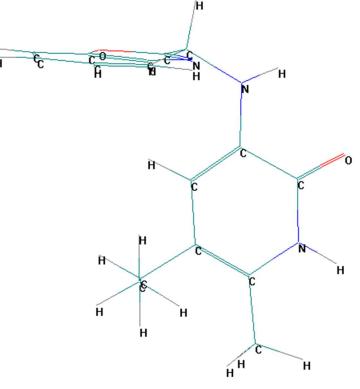
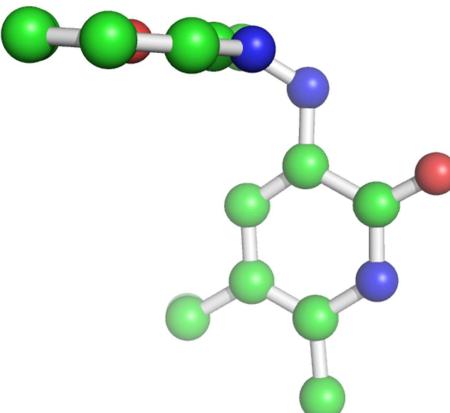
A

N

Lampiran 1. Nama 26 senyawa turunan 3-aminopiridin-2(1)-on

No	Nama Senyawa
1	3-(aminomethyl)phtahlimide-5-ethyl-6-methylpyridin-2(1H)-one
2	3-{[(benzoxazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
3	3-{[(benzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
4	3-[(4',5',6',7'-tetrahydrobenzoxazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
5	3-{{quinoline-3'yl} methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
6	3-{[(1',3'-benzothiazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
7	3-{[(2'naphtyl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
8	3-{[(benzothiophene-2'-yl)methyl] amino}-5-ethyl-6-methylpyridin-2(1H)-one
9	3-{[(quinoline-2'yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
10	3-{[(1'-benzopyran-4'-one-3'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
11	3-{[(3',4'-diazobenzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
12	3-naphthalene-5-ethyl-6-methylpyridin-2(1H)-one
13	3-{[(5'-phenyl-oxazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
14	3-{[(4'-quinoxolone-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
15	3-{[(1',3'-naftoxazol-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
16	3-{[(indol-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
17	3-benzylamino-5-ethyl-6-methylpyridin-2(1H)-one
18	3-{[(quinazolin-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
19	3-[(1',2',3',4'-tetrahydronaphthalene)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
20	3-{[(1',4'-naftoxazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
21	3-{[(pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
22	3-{[(indol-3'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
23	3-{[(furan-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
24	3-{[(5'-azabenzofuran-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
25	3-{[(6'-azabenzofuran-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one
26	3-{[(pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

Lampiran 2. Struktur hasil optimasi menggunakan program HyperChem

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
1	3-(aminomethyl)phtahlimide-5-ethyl-6-methylpyridin-2(1H)-one			-92053.843
2	3-{{[benzoxazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-81699.794

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
3	3-{[(benzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-80201.187
4	3-{[(4',5',6',7'-tetrahydrobenzoxazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-82997.401

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
5	3-{{[quinoline-3'yl]methyl}amino}-5-ethyl-6-methylpyridin-2(1H)-one			-80875.359
6	3-{{[(1',3'-benzothiazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-78816.119

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
7	3-{[(2'naphtyl) methyl]amino}-5-ethyl-6- methylpyridin-2(1H)-one			-79377.598
8	3-{[(benzothiophene-2'- yl)methyl] amino}-5-ethyl- 6-methylpyridin-2(1H)-one			-77318.849

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
9	3-{{[quinoline-2'yl] methyl}amino}-5-ethyl-6-methylpyridin-2(1H)-one			-80873.708
10	3-{{[(1'-benzopyran-4'-one-3'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-90550.798

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
11	3-{[(3',4'-diazobenzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-83192.940
12	3-naphthalene-5- ethyl-6-methylpyridin-2(1H)-one			-79375.318

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
13	3-{[(5'-phenyl-oxazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-88236.541
14	3-{[(4'-quinozolone-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-89771.061

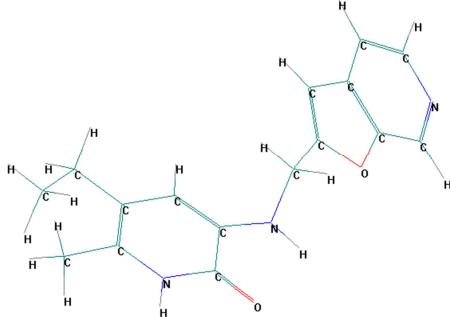
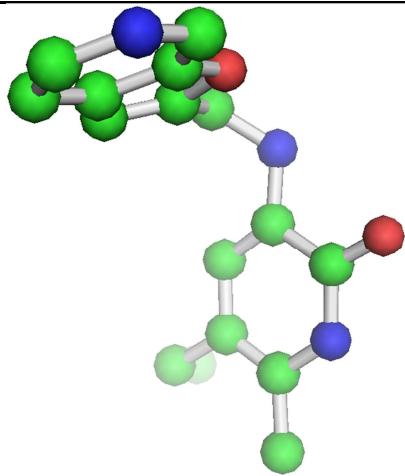
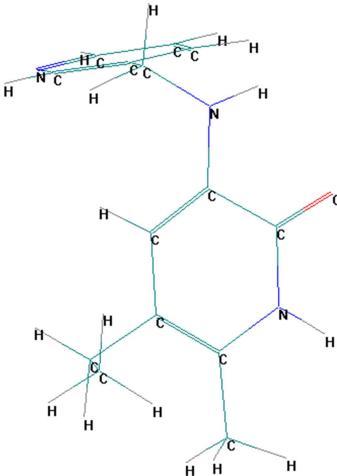
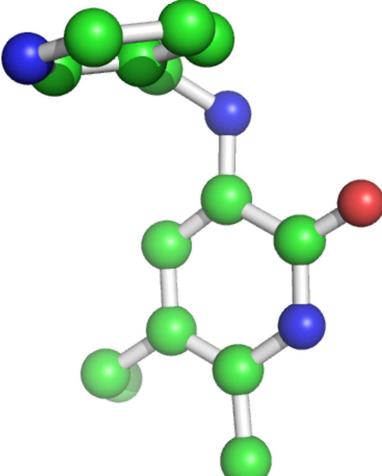
No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
15	3-{[(1',3'-naftoxazol-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-94139.974
16	3-{[(indol-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-77918.154

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
17	3-benzylamino-5-ethyl-6-methylpyridin-2(1H)-one			-66937.733
18	3-{{[(quinazolin-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-79415.161

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
19	3-[{(1',2',3',4'-tetrahydronaphthalene)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-80672.689
20	3-{[(1',4'-naftoxazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-94138.225

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
21	3-{{[pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-68435.573
22	3-{{[(indol-3'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-77916.125

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
23	3-{[(furan-2'-yl) methyl]amino}-5-ethyl-6- methylpyridin-2(1H)-one			-67761.234
24	3-{[(5'-azabenzofuran-2'- yl) methyl]amino}-5-ethyl-6- methylpyridin-2(1H)-one			-81699.309

No	Nama Senyawa	Gambar struktur optimasi 2D	Gambar struktur optimasi 3D	$\Delta G$ (kkal/mol)
25	3-{[(6'-azabenzofuran-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-77875.923
26	3-{[(pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one			-68436.844

Lampiran 3. Pola interaksi 26 senyawa turunan 3-aminopiridin-2(1H)-on dengan asam amino sesuai jenis interaksinya pada enzim HIV-1

No.	Turunan	Interaksi Hidrofobik	Interaksi Hidrogen
1.	Turunan 1	Asn265, Gln 269, Trp269, Met230, Tyr232, Ile94, His94, Ile382, dan Lys350.	Glu378
2.	Turunan 2	Pro225, Pro236, His235, Leu234, Tyr318, Lys103, Tyr188, Tyr181, Leu100, Lys101, Val106, Phe227, dan Lys223.	
3.	Turunan 3	Lys22, Lys20, Asn57, Val21, Gln23, Pro140, Thr131 Gly141, Pro52, Arg143, Tyr56, dan Pro55.	
4.	Turunan 4	Pro225, Ser105, Pro236, Val106, Lys223, Phe227, His235, Tyr188, Leu234, Leu100, Tyr181, Tyr318, Lys103, dan Lys104.	
5.	Turunan 5	Lys22, Val21, Lys20, Pro55, Tyr56, Arg143, Pro 52, Pro140, dan Asn57.	Gly141, Gln23, Thr131.
6.	Turunan 6	Pro236, Val106, Tyr318, Lys101, Tyr181, Tyr188, Leu234, Trp229, Pro95, Leu100, His235, dan Lys103.	
7.	Turunan 7	Pro236, Leu100, Tyr181, Val179, Glu190, Tyr188, Tyr318, Leu234, Pro225, His235, Phe227, Ser105, Lys104, dan Val106.	Lys103
8.	Turunan 8	Pro225, His235, Ser105, Pro236, Lys104, Leu100, Val179, Gly190, Tyr181, Tyr188, Tyr318, Phe227, Val106, dan Leu234.	Lys103
9.	Turunan 9	Leu100, Tyr181, Trp229, Tyr188, Val108, Leu234, Phe227, Val106, Tyr318, dan Lys103	Lys101
10.	Turunan 10	Asn265, Gln 269, Ser268, Glu378, Val381, Tyr232, , His96, dan Ile382	Lys350, Ile94
11.	Turunan 11	Pro236, His235, Ser105, Lys223, Phe227, Val106, Leu100, Lys101, Tyr181, Tyr318, Tyr188, dan Leu234.	Lys103, Lys104
12.	Turunan 12	Pro236, Lys103, Val106, Leu234, Phe227, Tyr181, Tyr188, Leu100, Lys101, dan Tyr318	
13.	Turunan 13	Pro236, Pro225, Val106, His235, Phe227, Tyr188, Leu234, Pro95, Trp229, Leu100, Tyr181, Lys103, dan Tyr318.	Lys101
14.	Turunan 14	Lys311, Arg307, Leu310, Gln242, Thr240, Pro243, Glu233, Pro313, dan Leu246.	Tyr271, Ile244, Val241.
15.	Turunan 15	Lys311, Arg307, Leu310, Gln242, Thr240, Pro243, Glu233, dan Pro313.	Tyr271, Ile244, Val241.

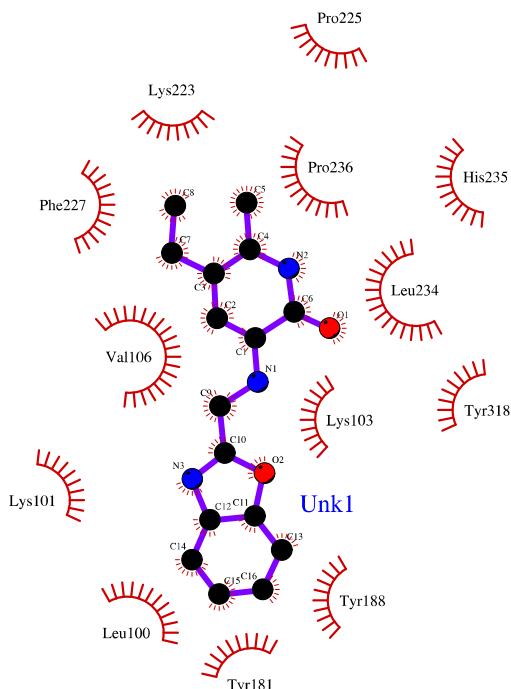
---

16. Turunan 16	Pro236, Tyr318, His235, Pro225,Lys223, Phe227, Ser105, Val106, Leu234, Tyr188, Tyr181, Lys101, Val179, dan Lys104.	Lys103
17. Turunan 17	Tyr318, His235, Phe227, Leu234, Trp229, Tyr188, Leu100, Tyr181, Lys103, dan Val106.	Lys101
18. Turunan 18	Pro236, Lys103, Leu100, Pro95, Trp229, Tyr188, Tyr181, Leu234, Val106, Lys101, His235, dan Tyr318.	
19. Turunan 19	Asn265, Ser268, Glu378, Lys353, Tyr339, Gly352, His96, Ile94, Trp266, dan Ile382.	Lys350
20. Turunan 20	Pro236, Ser105, Lys104, Val106, Leu234, Lys103, Tyr318, Val179, Leu100, Tyr181, His235, Phe227, Lys223, dan Pro225.	Lys101
21. Turunan 21	Pro236, Val106, Phe227, Leu234, Tyr318, Tyr188, Leu100, Tyr181, Val179, His235, dan Lys104.	Lys103
22. Turunan 22	Pro236, Lys103, Tyr188, Tyr181, Leu100, Pro95, Trp229, Leu234, Val106, His235 Phe227, dan Tyr318.	Lys101
23. Turunan 23	Val106, Phe227, Lys103, Leu100, Trp229, Leu234, Tyr181, Tyr188, dan Tyr 318.	Lys101
24. Turunan 24	Val241, Lys311, Arg307, Pro243, Leu246, dan Leu310.	Tyr271, Ile244
25. Turunan 25	Pro236, His235, Phe227, Leu234, Lys104, Tyr318, Val106, Leu100, Val189, Val179, Glu190, Tyr181, dan Pro225.	Lys103, Tyr188
26. Turunan 26	Lys22, Gln23, Asn57, Val21, Lys20, Tyr56, Pro55, Pro140, Gly141, dan Arg143.	Thr131

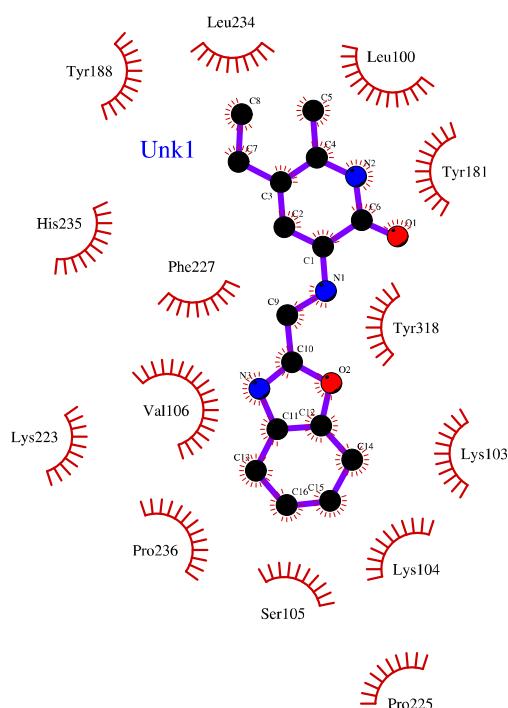
---

Lampiran 4. Pola interaksi 17 senyawa turunan senyawa 3-aminopiridin-2(1H)-on menempati daerah penghambatan yang sama dengan ligan pada daerah (a) enzim RT HIV-1.

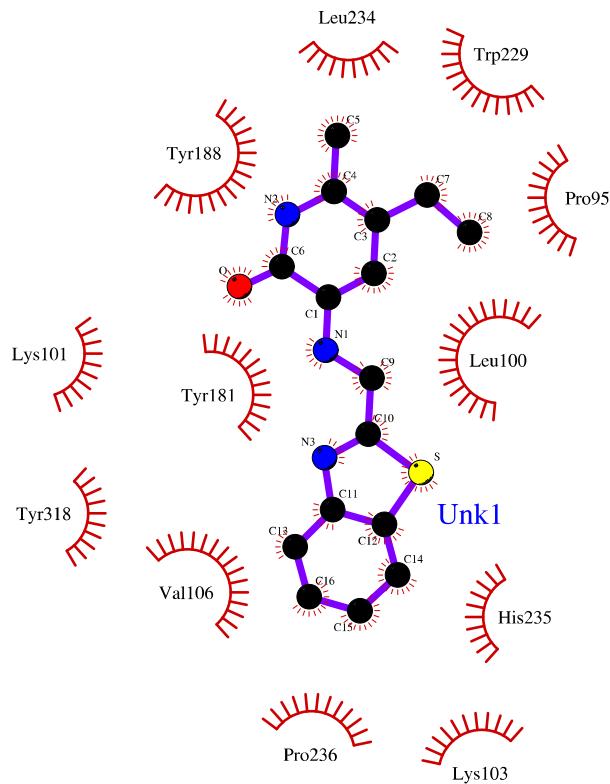
2. 3-{[(benzoxazole-2'-yl)methyl]amino}5-ethyl-6-methylpyridin2(1H)-one



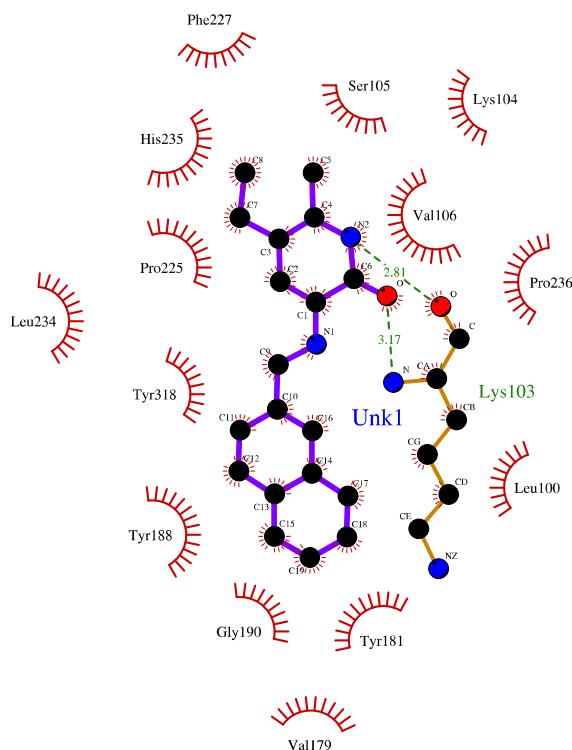
4. 3-{[(4',5',6',7'-tetrahydrobenzoxazole-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



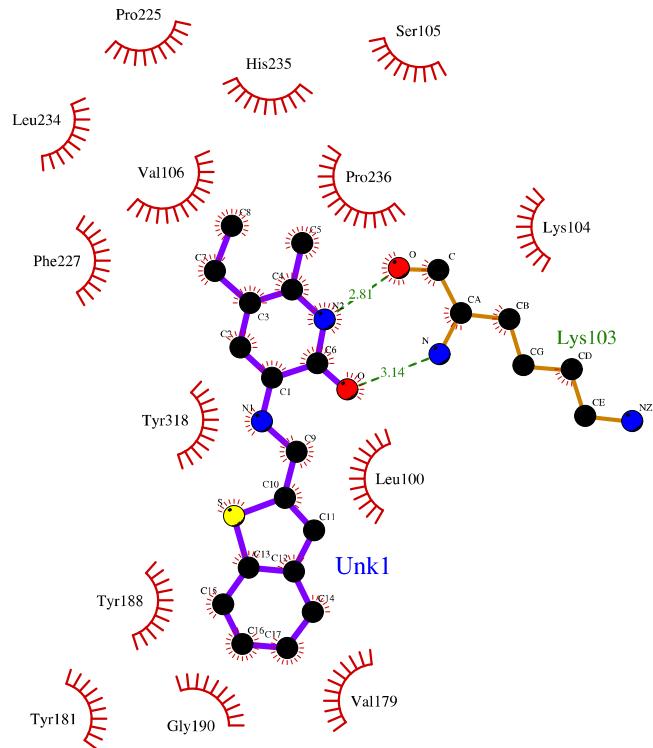
6. 3-{[(1',3'-benzothiazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



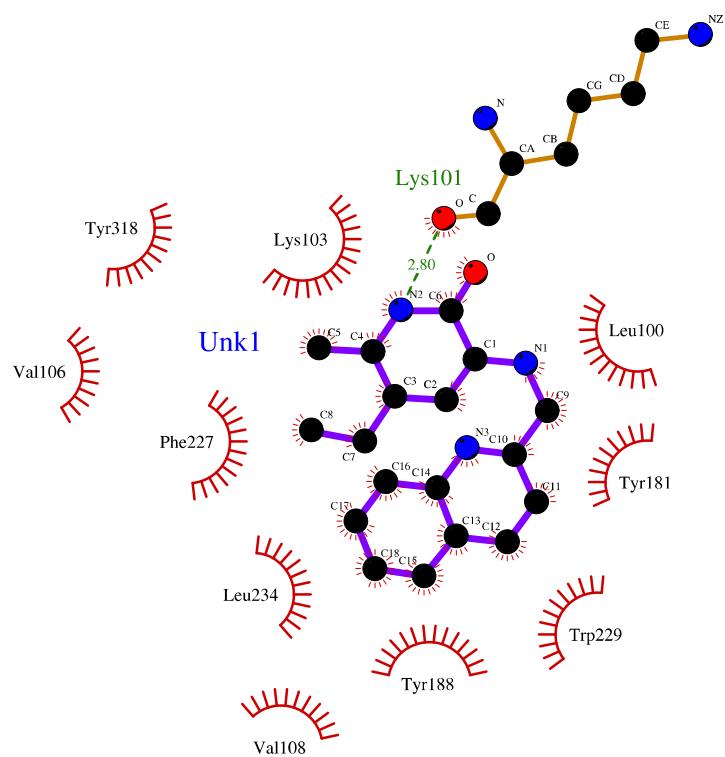
7. 3-{[(2'naphthyl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



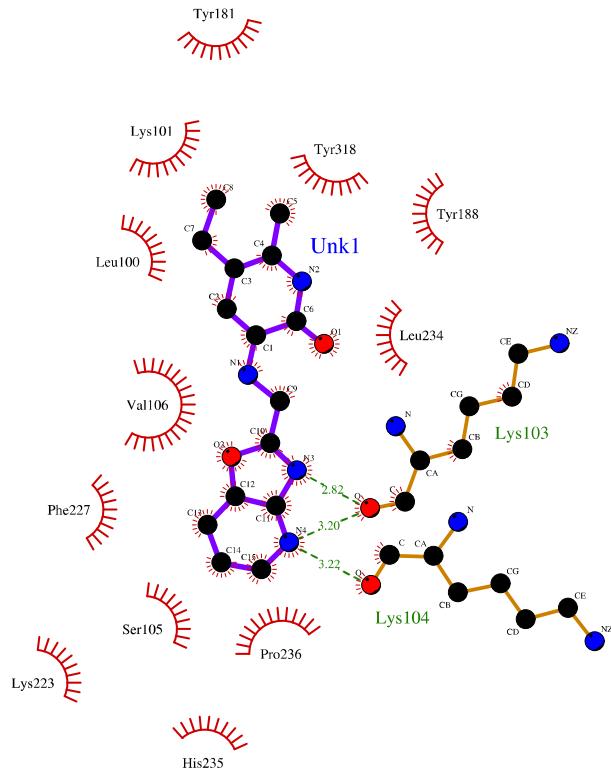
8. 3-{[(benzothiophene-2'-yl)methyl] amino}-5-ethyl-6-methylpyridin-2(1H)-one



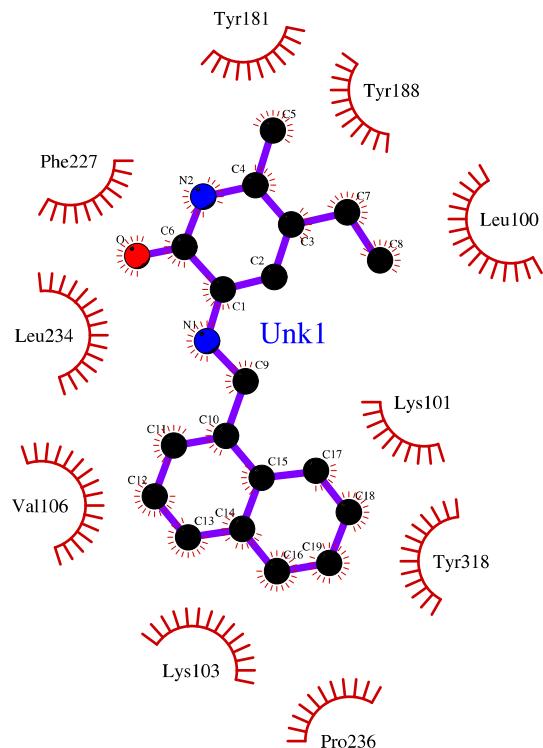
9. 3-{[quinoline-2'yl] methyl}amino}-5-ethyl-6-methylpyridin-2(1H)-one



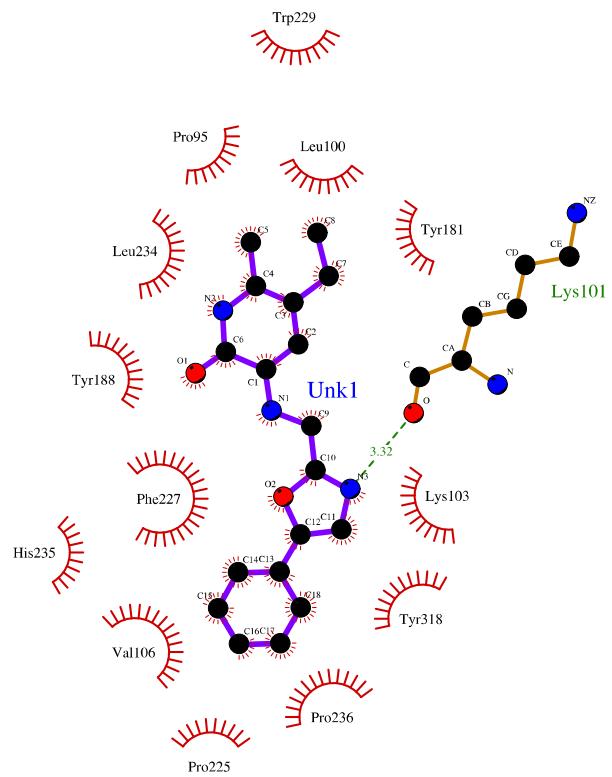
11. 3-{[(3',4'-diazobenzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



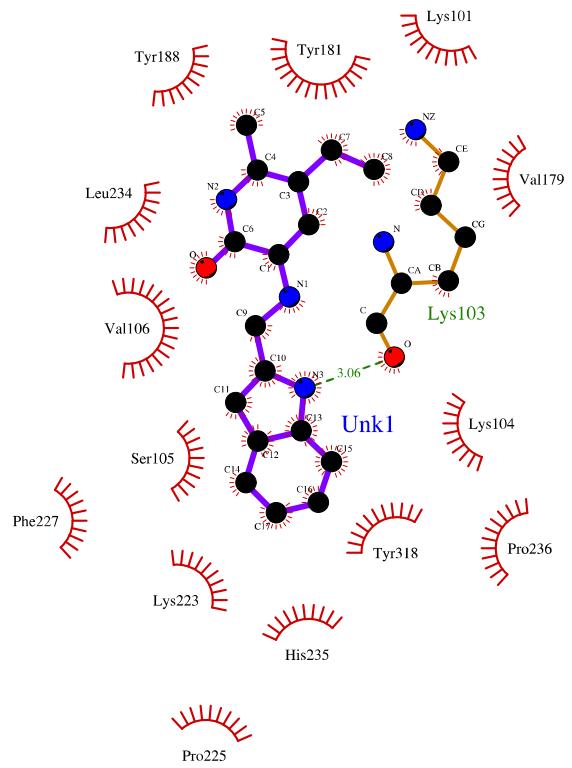
12. 3-naphthalene-5- ethyl-6-methylpyridin-2(1H)-one



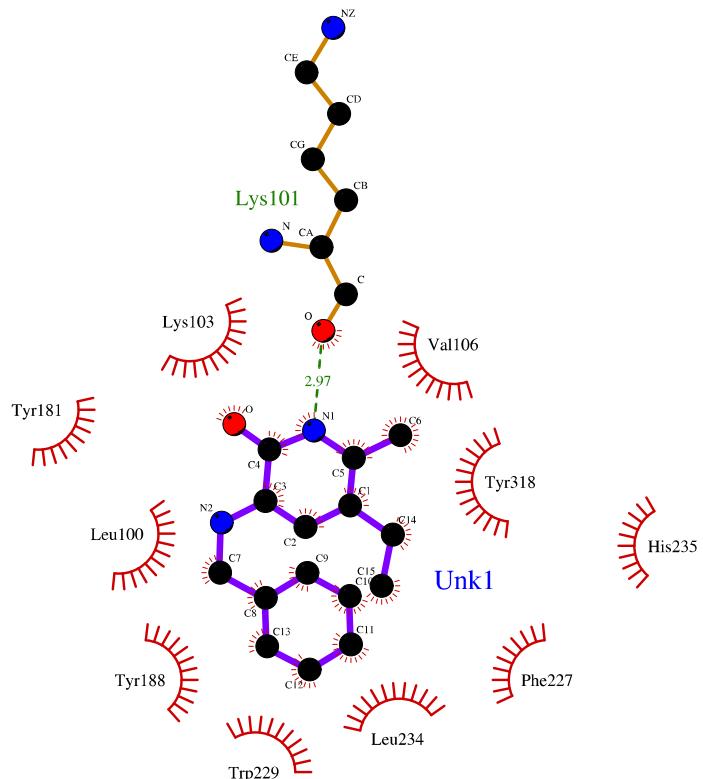
13. 3-[(5'-phenyl-oxazole-2'-yl) methyl]amino]-5-ethyl-6-methylpyridin-2(1H)-one



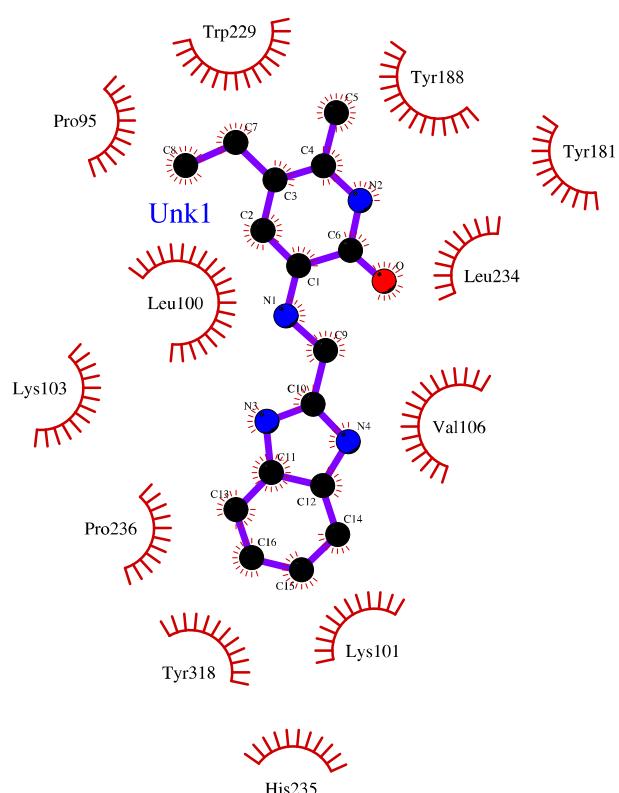
16. 3-[(indol-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



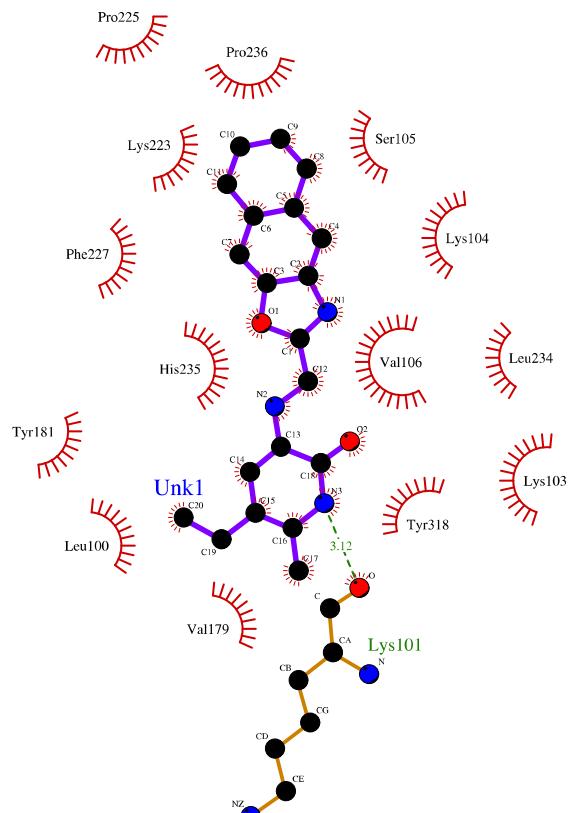
17. 3-benzylamino-5-ethyl-6-methylpyridin-2(1H)-one



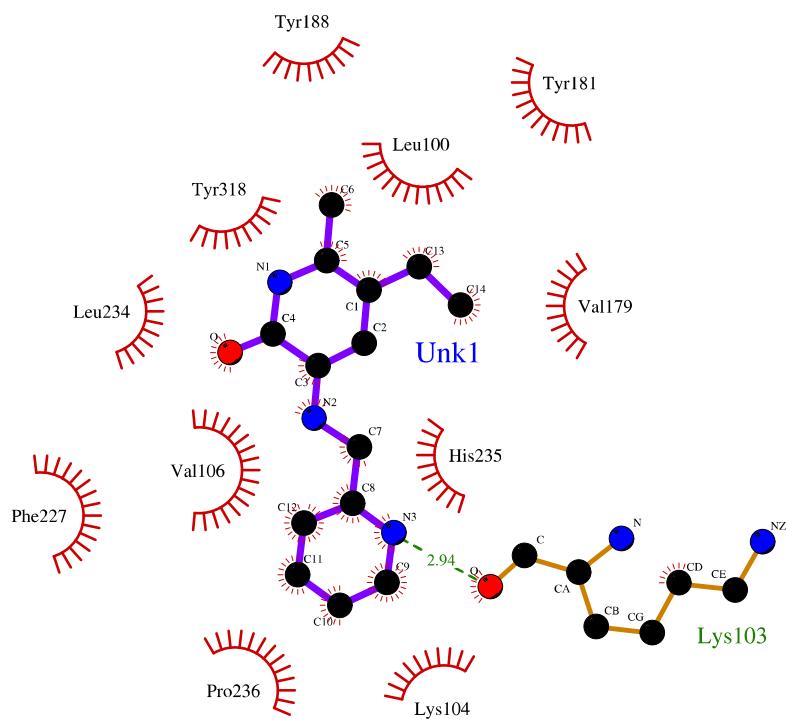
18. 3-{{[(quinazolin-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



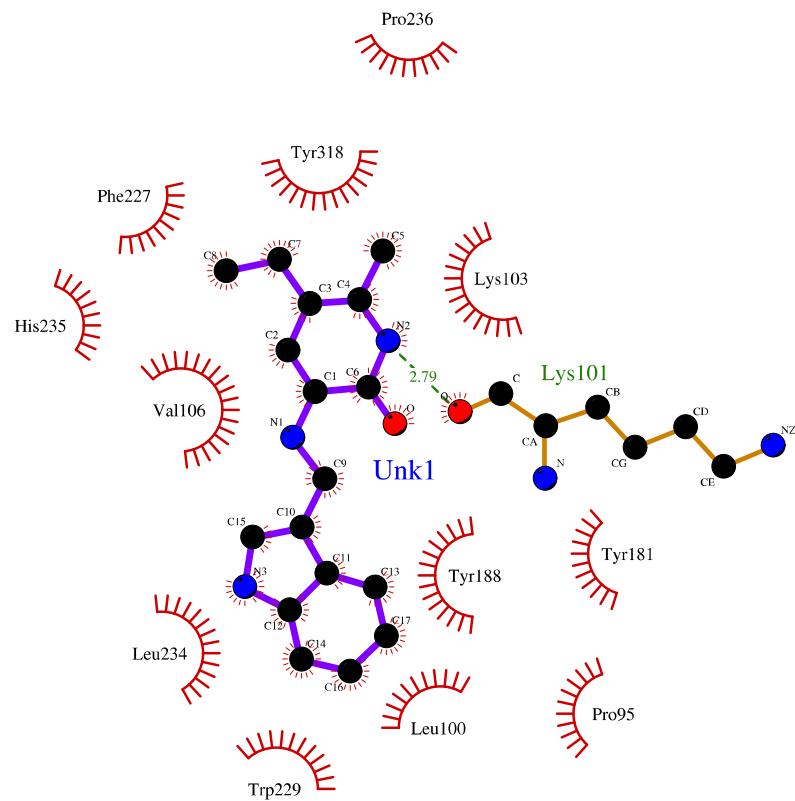
20. 3-{[(1',4'-naftoxazole-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



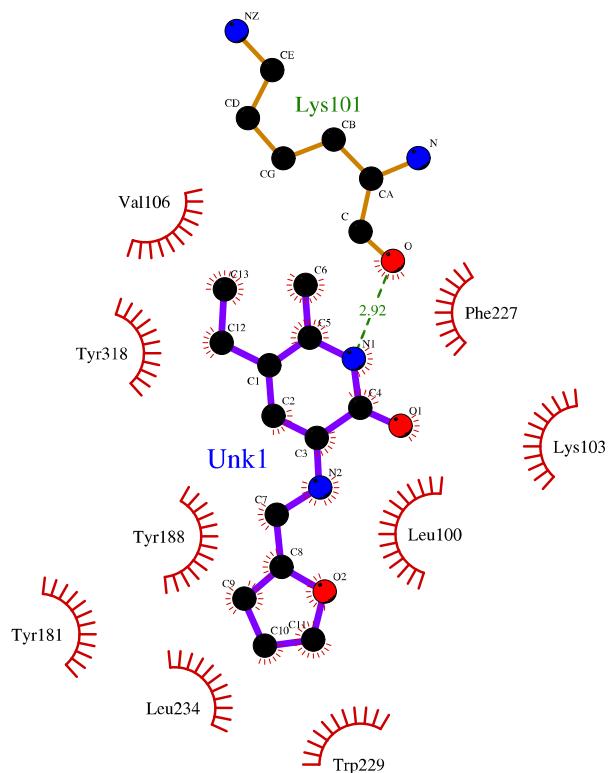
21. 3-{[(pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



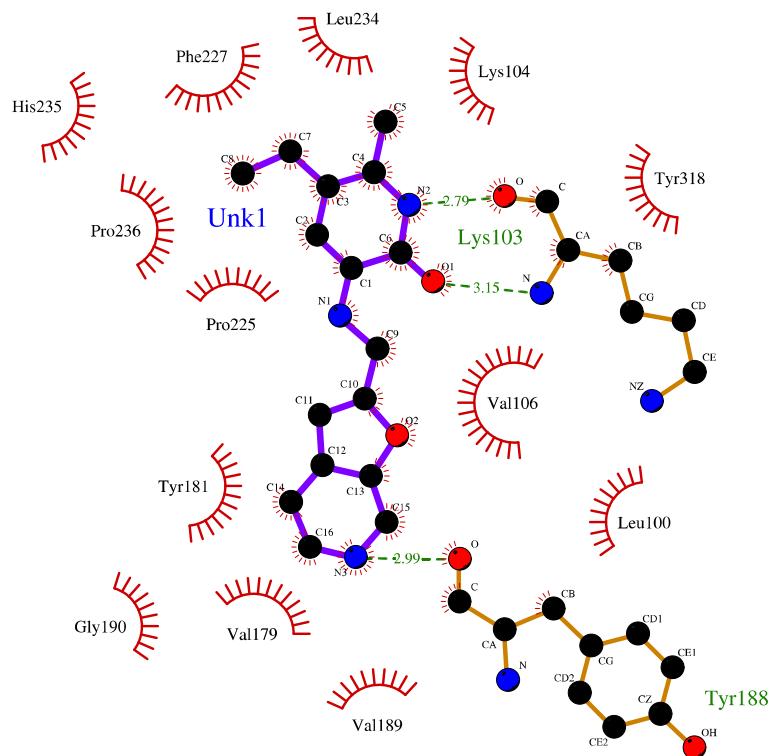
22. 3-{[(indol-3'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



23. 3-{[(furan-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

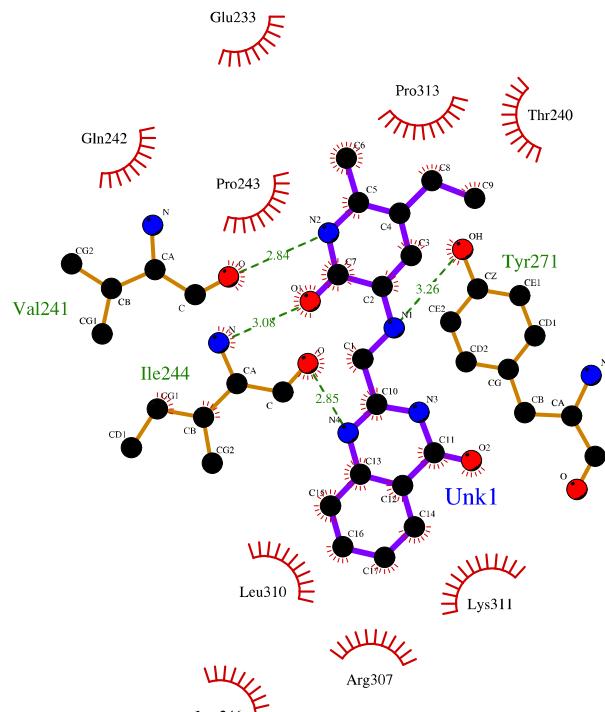


25. 3-{[(6'-azabenzofuran-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

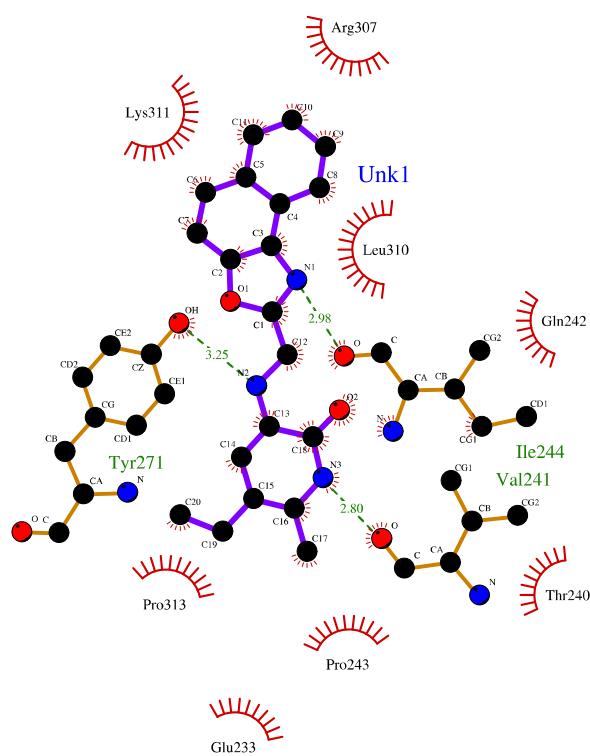


Lampiran 5. Pola interaksi 3 senyawa turunan 3-aminopiridin-2(1H)-on terhadap *binding site* RT HIV-1 pada daerah (b).

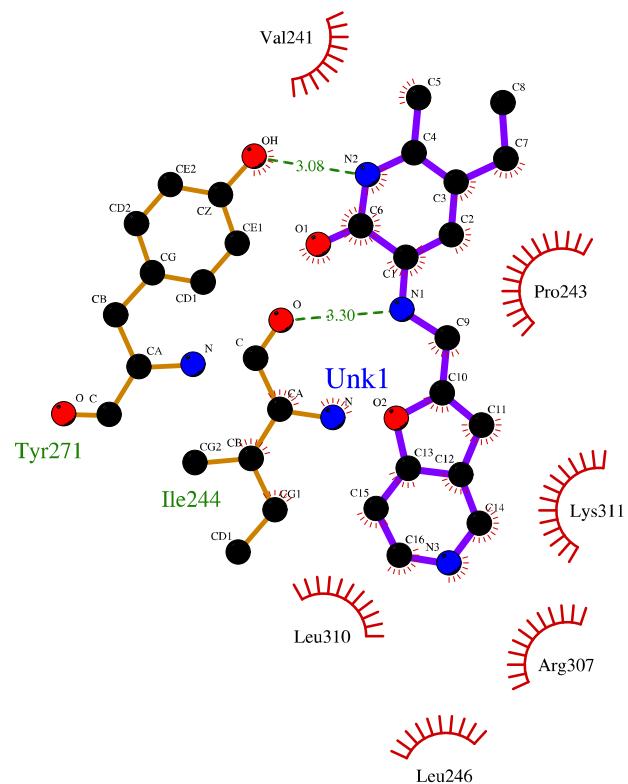
14. 3-{[(4'-quinoxolone-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



15. 3-{[(1',3'-naftoxazol-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

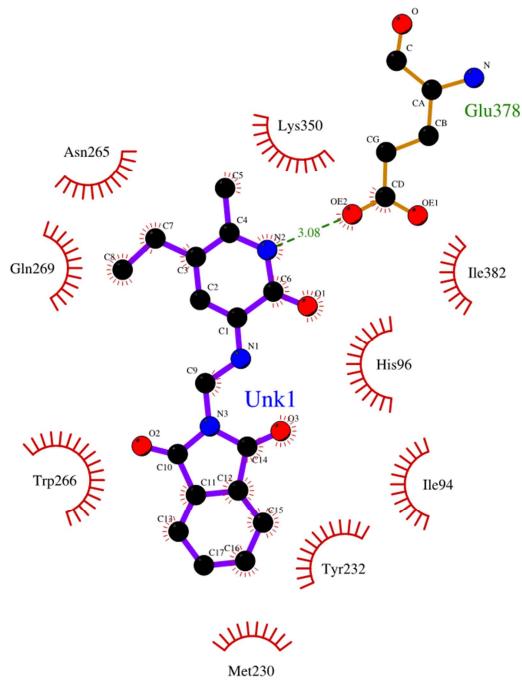


24. 3-{[(5'-azabenzofuran-2'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

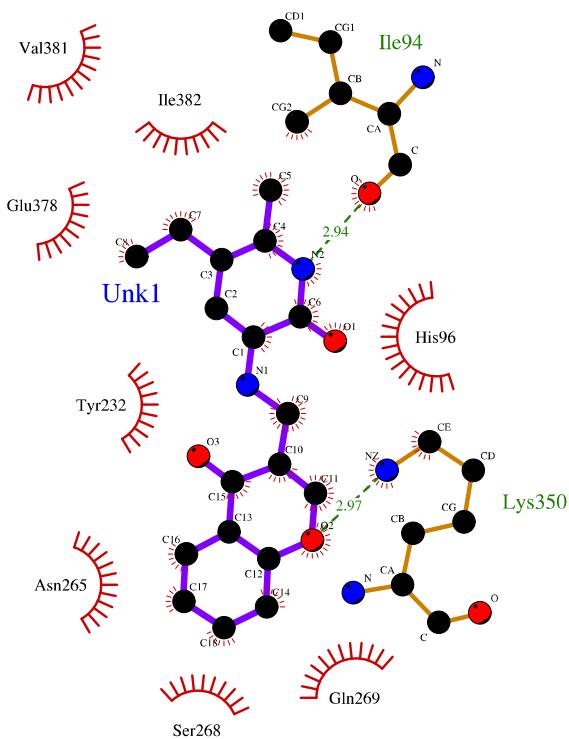


Lampiran 6. Pola interaksi 3 senyawa turunan 3-aminopiridin-2(1H)-on terhadap *binding site* RT HIV-1 pada daerah (c).

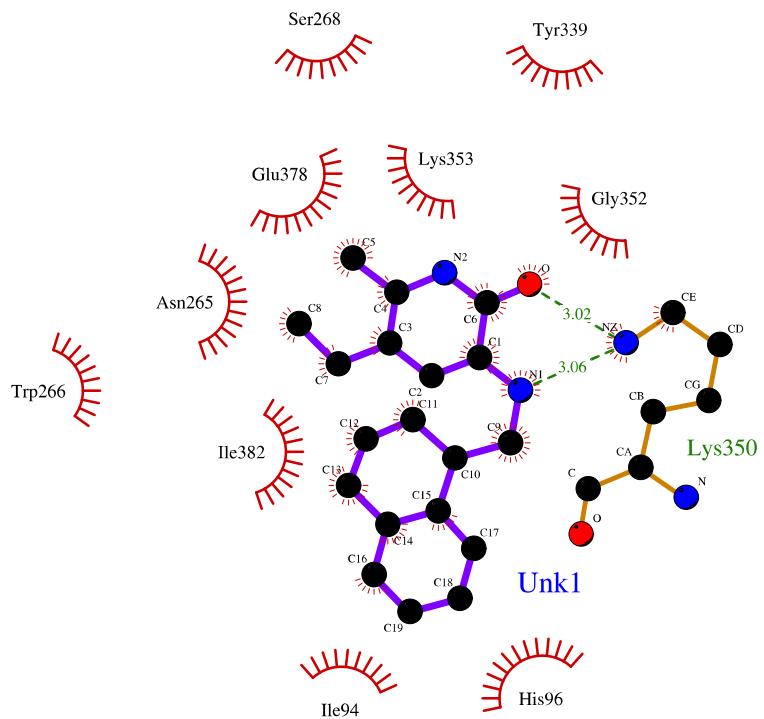
1. 3-(aminomethyl)phtahlimide-5-ethyl-6-methylpyridin-2(1H)-one



10. 3-{[(1'-benzopyran-4'-one-3'-yl) methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

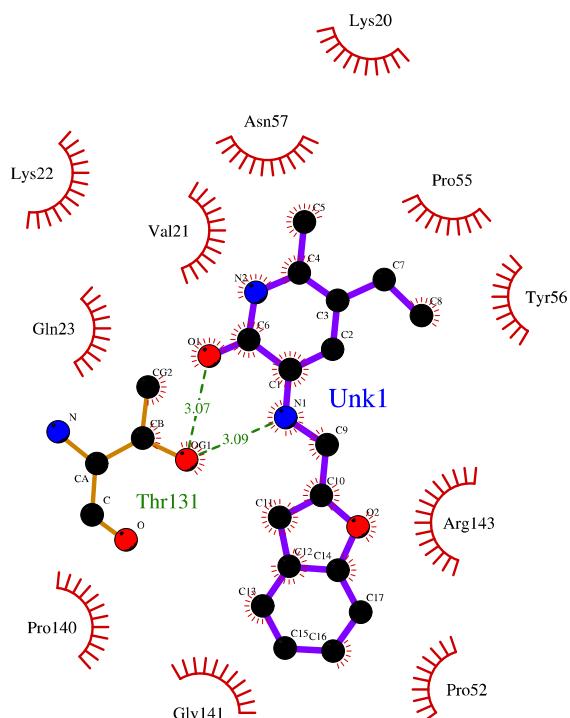


19. 3-[(1',2',3',4'-tetrahydronaphthalene)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

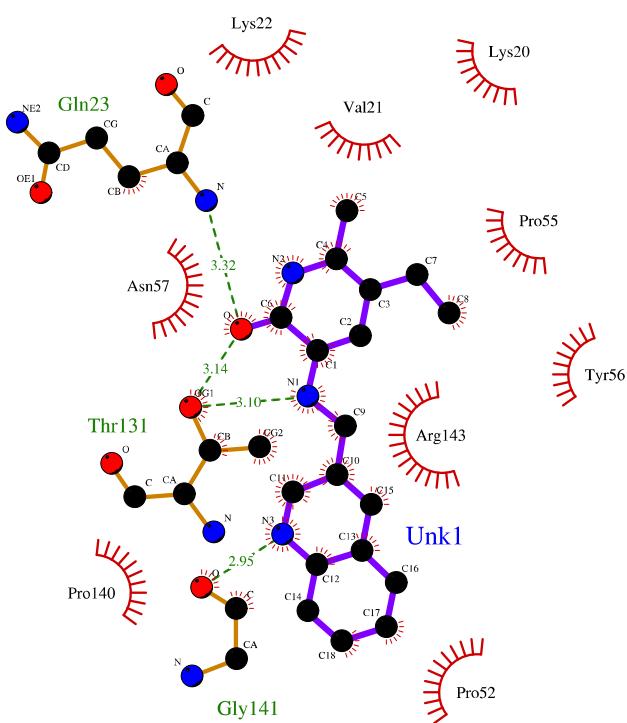


Lampiran 7. Pola interaksi 3 senyawa turunan 3-aminopiridin-2(1H)-on terhadap *binding site* RT HIV-1 pada daerah (d).

3. 3-{[(benzofuran-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one



5. 3-{[quinoline-3'yl] methyl}amino}-5-ethyl-6-methylpyridin-2(1H)-one



## 26. 3-{[(pyridine-2'-yl)methyl]amino}-5-ethyl-6-methylpyridin-2(1H)-one

