

Design, Synthesis and Evaluation of Novel 1-(Substituted Acetyl)-4-(10-Bromo-8-Chloro-5,6-Dihydro-11H-Benzo[5,6]Cyclohepta[1,2-B]Pyridine-11-Ylidene) Piperidines as Antitumor Agents and Farnesyl Protein Transferase Inhibitors

P. S. GATNE*, C. L. VISWANATHAN*, PREMLATA K. AMBRE AND AARTI JUVEKAR¹

Department of Pharmaceutical Chemistry, Bombay College of Pharmacy, Kalina, Santacruz, Mumbai-400 098,

¹Anticancer Drug Screening Facility (ACDSF), Tata Memorial Centre, Advanced Center for Treatment Research and Education in Cancer (ACTREC), Kharghar, Navi Mumbai-410 210, India

Gatne, *et al.*: Dihydro-11H-Benzo[5,6]Cyclohepta[1,2-B]Pyridine-11-Ylidene)Piperidines as Antitumor Agents

Eight novel 1-(substituted acetyl)-4-(10-bromo-8-chloro-5,6-dihydro-11H-benzo[5,6] cyclohepta [1,2-b] pyridine-11-ylidene)piperidines were designed by incorporating zinc binding groups to enhance activity. The designed molecules were synthesized and were evaluated for antitumor activity *in vitro* in five cell lines and for farnesyl protein transferase inhibition. Test compounds (6a-h) exhibited antitumor activity in most of the cell lines but were less potent than adriamycin. Compound 6e was most active with IC₅₀ values of <15 µM in two cell lines tested. Test compounds also exhibited potent FPT inhibitory activity and 6c was most potent with IC₅₀ value of <30 µM.

Key words: Anticancer agents, benzocycloheptapyridines, farnesyl protein transferase inhibitors

*Address for correspondence

E-mail: parag507@gmail.com

An important therapeutic approach in cancer is to inhibit farnesylation of ras and this can be achieved by inhibition of farnesyl protein transferase (FPT) enzyme using inhibitors (FTIs)^[1]. FTIs belonging to various structural classes were developed and among these, lonafarnib[®] (benzocycloheptapyridines)^[2], tipifarnib[®] (4,6-disubstituted-quinoline-2-ones)^[3] and BMS-241662 (1,3,4-trisubstituted-2,3,4,5-tetrahydro-1,4-benzodiazepines)^[4] are potent molecules and are currently undergoing phase II and III clinical trials.

Among benzocycloheptapyridine class of FTIs, the tricyclic nucleus, a bromine at C₁₀ position on the ring, a piperidine ring at C₁₁ position linked to heteroaryl systems by a -COCH₂ linker are important structural parameters for activity. It is also reported that the incorporation of groups with zinc binding ability at 2-position of the piperidine ring can enhance activity (structure I, fig. 1)^[5].

Hence in the present work, it is proposed to develop a series of novel benzocycloheptapyridines with structure II (fig. 1) where *R* is a group of heteroaryl ring systems. It is also proposed that the group *R* in structure II would occupy the zinc binding region in FPT and thereby exhibit potent activity. Hence for designing of novel molecules, the 'Flexible Alignment' tool of Molecular Operating Environment 2006.08 (MOE, Chemical Computing group, USA) software was used on a Windows platform with a Pentium IV Dual Core processor computer (2.9 GHz and 1 GB RAM).

Conformers of I, lonafarnib[®] and test molecules (6a-h) (general structure II) were generated by using dynamic simulations at the temperature of 310° and the sampling time was 5×10⁻⁴ s and the lowest energy conformations were identified for each molecule. Keeping I as template, lonafarnib and 6a-h were superimposed with similarity terms being Aromaticity, Hydrophobe, LogP (o/w) and Hydrogen bond acceptor. Similarity index (F); Average strain energy (U) and Alignment score (S) were measured and are given in Table 1. Lower scores indicate greater similarity and better alignment. Alignment of I with lonafarnib (fig. 2a) and with one of the test compounds (6e) is given in fig. 2b. From fig. 2b, it is clear that alignment of 6e is highly significant compared to that of lonafarnib.

Loratadine[™] was obtained as a gift sample from

Themis laboratories, Thane, India and was used as a starting material for the synthesis of 6a-h. ³[H] FPP, H-ras protein and FPT required for the assay were purchased from Sigma Aldrich (USA). Desloratadine was obtained as a gift sample from Glenmark Pharmaceuticals, Mahape, Navi Mumbai, India.

10-Bromodesloratadine (3) was prepared from loratadine (1) by nitration using conc. sulfuric acid and potassium nitrate at -10° for 30 min to get a mixture of two nitro isomers (9-nitro- and 7-nitro-loratadine). The nitro group in the mixed isomers was then reduced to amine using stannous chloride dihydrate in ethyl acetate at room temperature.

TABLE 1: SCORES OBTAINED FOR FLEXIBLE ALIGNMENT STUDY

Compound	Scores		
	F	S	U
6a	179.31	293.51	114.20
6b	176.94	298.13	126.61
6c	170.74	272.01	113.10
6d	172.10	286.50	109.86
6e	167.72	276.68	103.95
6f	165.33	273.55	108.22
6g	169.72	276.25	106.00
6h	174.30	281.84	101.42
Lonafarnib	164.95	253.45	78.65

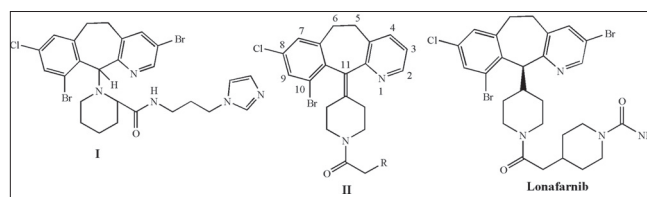


Fig. 1: Structures of I, II and lonafarnib
Molecules from benzocycloheptapyridine class of farnesyl protein transferase inhibitors which are currently under investigation

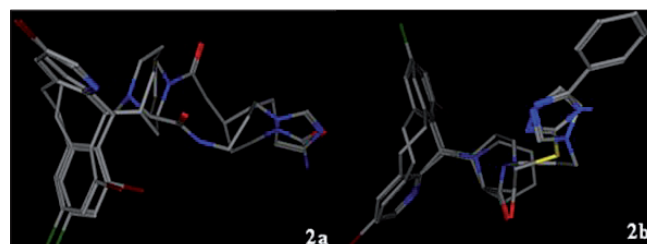


Fig. 2: Flexible alignment of lonafarnib with compound I and test compounds (6e).

Alignment of I with lonafarnib (2a) and with one of the test compounds (6e) is given in fig. 2b. From 2b, it is clear that alignment of 6e is highly significant compared to that of lonafarnib

The mixed amines formed were then brominated using bromine in acetic acid at 15-20° to achieve bromination at the C₁₀ position on ring. Diazotization of amine function with sodium nitrite and concentrated HCl at 0° followed by treatment with hypophosphorous acid at 5° gave 10-bromoloratadine (2) as a single isomer. 10-bromoloratadine was decarboethoxylated using sodium hydroxide in methanol at reflux to get 10-bromodesloratadine. Further reaction with chloroacetyl chloride gave an intermediate (4), which was then condensed with various substituted amines or thiols (5a-h) in dimethyl formamide (DMF) in presence of base like potassium carbonate or sodium hydride to get test compounds 6a-h (Scheme 1).

Substituted amines/thiols 5d-5f and 5h were synthesized as per literature methods^[6,7], while 4-amino-5-phenyl-3-thiol-1,2,4-triazole (5g) was synthesized from 5-phenyl-1,3,4-oxadiazole-2-thiol by reacting it with 40% methyl amine solution under microwave conditions. Amines 5a-5c were obtained as gift samples from RPG Lifesciences, Pawane, Navi Mumbai, India.

The yield, mp, IR and NMR and mass spectral characteristics of 10-bromoloratadine, 10-bromodesloratadine and test compounds 6a-h are given as follows, 10-bromoloratadine (2), Yield: 65%;

mp: 174-176°; IR (KBr): 3059 (C-H stretch, Ar), 2972,1429 (C-H stretch, aliph), 1695 (C=O stretch, ester), 1224 (C-O stretch, acetate), 769 (C-Cl stretch), 525 (C-Br stretch); NMR (CDCl₃): 8.47 (s, 1H, Ar), 7.46 (s, 1H, Ar), 7.37 (d, 1H, Ar), 7.20 (s, 1H, Ar), 7.1 (m, 1H, Ar), 4.15 (q, 2H, Aliph), 3.84 (s, 2H, Aliph), 3.14-3.50 (m, 4H, Aliph), 2.8 (m, 2H, Aliph), 2.6 (m, 1H, Aliph), 2.3-2.4 (m, 2H, Aliph), 2.0 (m, 1H, Aliph), 1.25 (t, 3H, Aliph); GC-MS (ES)- 462.5.

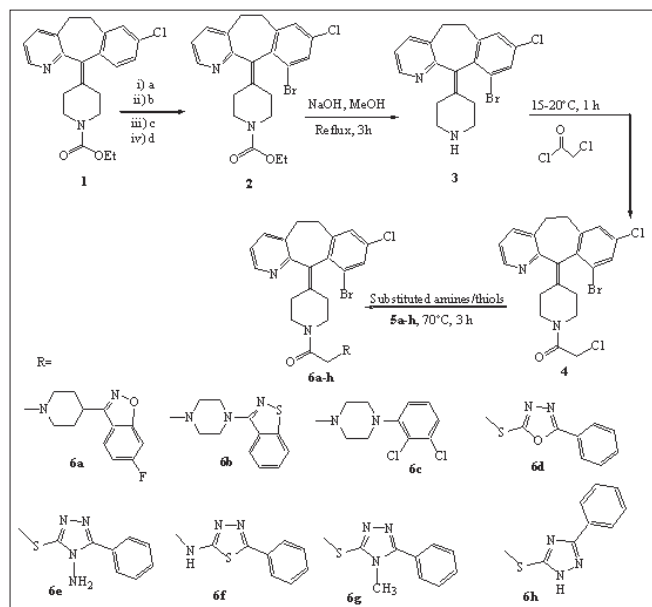
10-bromodesloratadine (3), Yield: 65%; mp: 114-116°; IR (KBr): 3508-3385 (N-H stretch, secondary aliphatic amines), 3059 (C-H stretch, Ar), 2928, 1429 (C-H stretch, Aliph), 1103 (C-N stretch, aliphatic amine), 798 (C-Cl stretch), 525 (C-Br stretch); LC-MS (ES)- 388.98, 390.96, 392.95, 393.97.

6a, Yield: 80%; mp: 116-118°; IR (KBr): 3061 (C-H stretch, Ar), 2928,1446 (C-H stretch, Aliph), 1643 (C=O, amide stretch), 1300 (C-N stretch, tertiary amine), 1000 (C-O stretch, ether), 787 (C-Br stretch), 675 (C-Cl stretch); ¹H-NMR (CDCl₃): 8.5 (s, 1H, Ar), 7.7 (s, 1H, Ar), 7.5 (s, 1H, Ar), 7.37 (d, 1H, Ar), 7.26 (m, 2H, Ar), 7.1 (m, 2H, Ar), 3.9 (s, 2H, Aliph), 3.2-3.6 (m, 6H, Aliph), 3.1 (d, 3H, Aliph), 2.2-2.9 (m, 7H, Aliph), 2.1 (m, 5H, Aliph).

6b, Yield: 70%; mp: 110-112°; IR (KBr): 3059 (C-H stretch, Ar), 2903, 1423 (C-H stretch, Aliph), 1641 (C=O stretch, amide), 1350 (C-N stretch, tertiary amine), 787 (C-Cl), 525 (C-Br stretch); ¹H-NMR (CDCl₃): 8.5 (s, 1H, Ar), 7.9 (m, 2H, Ar), 7.4 (d, 2H, Ar), 7.36 (m, 2H, Ar), 7.26 (m, 2H, Ar), 3.85 (m, 2H, Aliph), 3.2-3.6 (10H, Aliph), 2.9 (m, 7H, Aliph), 2.4 (m, 2H, Aliph), 2.1 (m, 1H, Aliph); LC-MS (ES)- 647.99, 649.8, 650.04 (MH⁺).

6c, Yield: 70%; mp: 104-106°; IR (KBr): 3059 (C-H stretch, Ar), 2920, 1448 (C-H stretch, Aliph), 1641 (C=O stretch, amide), 1275 (C-N stretch, tertiary amine), 787 (C-Br stretch), 711 (C-Cl stretch); ¹H-NMR (CDCl₃): 8.5 (s, 1H, Ar), 7.5 (s, 1H, Ar), 7.43 (d, 1H, Ar), 7.2(m, 5H, Ar), 4.1 (m,1H, Aliph), 3.9(m, 1H, Aliph), 3.4 (m, 6H, Aliph), 3.1 (m, 4H, Aliph), 2.9 (m, 7H, Aliph), 2.5 (m, 2H, Aliph), 2.1 (m, 1H, Aliph).

6d, Yield: 60%; mp: 118-122°; IR (KBr): 3059 (C-H stretch, Ar), 2987, 1467 (C-H stretch, Aliph), 1643 (C=O stretch, amide), 1525(C=N stretch, imines), 1320 (C-N stretch, tertiary amine), 990 (C-O stretch, ethers),



Scheme 1: Synthesis of test compounds 6a-h
 Reagents and conditions (a) concentrated H₂SO₄, KNO₃, -5°, 30 min; (b) SnCl₄·2H₂O, RT, 1 h; (c) Br₂, AcOH, 15°, 2 h; (d) i) NaNO₂, concentrated HCl, 0°, 1 h ii) hypophosphorous acid, 5°, 2 h

764 (C-Br stretch), 673 (C-Cl stretch); ¹H-NMR (CDCl₃): 8.5 (d, 1H, Ar), 8.0 (d, 2H, Ar), 7.6 (m, 4H, Ar), 7.4(d, 1H, Ar), 7.35 (s, 1H, Ar), 7.15(m, 1H, Ar), 4.4 (s, 2H, Aliph), 4.10 (m, 1H, Aliph.), 3.8 (m, 1H, Aliph), 3.3-3.8 (m, 4H, Aliph), 2.8 (m, 3H, Aliph), 2.4 (m, 2H, Aliph), 2.1 (m, 1H, Aliph).

6e, Yield: 65%; mp: 154-157°; IR (KBr): 3314-3182 (N-H stretch for amines), 3061 (C-H stretch, aromatic), 2924, 1457 (C-H stretch, Aliph), 1590 (C=N stretch, imines), 1639 (C=O stretch, amide), 1545 (N-H bending vibrations), 771 (C-Cl stretch), 694 (C-Br stretch); ¹H-NMR (CDCl₃) 8.35 (d, 1H, Ar), 8.15 (d, 2H, Ar), 7.45 (m, 4H, Ar), 7.26 (s, 1H, Ar), 7.20 (m, 2H, Ar), 5.45 (s, 2H, NH₂, D₂O exchangeable), 4.1(s, 2H, Aliph), 3.8 (m, 2 H, Aliph), 3.30 (m, 4H, Aliph), 2.7 (m, 4H, Aliph), 2.07(m, 2H, Aliph).

6f, Yield: 50%; mp: 170-174°; IR (KBr): 3061 (C-H stretch, aromatic), 2924, 1457 (C-H stretch, Aliph), 1552 (C=N stretch), 1639 (C=O stretch, amide), 771 (C-Cl stretch), 694 (C-Br stretch); ¹H-NMR (CDCl₃): 8.75 (d, 1H, Ar), 7.63 (d, 2H, Ar), 7.57 (m, 4H, Ar), 7.10(s, 1H, Ar), 7.05 (m, 2H, Ar), 5.9 (1H, N-H) 3.9 (s,2H, Aliph), 3.4 (5 H, Aliph), 2.80 (m, 3H, Aliph), 2.4 (m, 2H, Aliph), 2.1 (m, 2H, Aliph).

6g, Yield: 65%; mp: 157-160°; IR (KBr): 3061 (C-H stretch, aromatic), 2924, 1457 (C-H, Aliph), 1712 (C=N stretch), 1639 (C=O, amide), 771 (C-Cl stretch), 694 (C-Br stretch); ¹H-NMR (CDCl₃): 8.5 (d, 1H, Ar), 8.15 (d, 2H, Ar), 7.45 (m, 4H, Ar), 7.26 (s, 1H, Ar), 7.20 (m, 2H, Ar), 4.1(s, 2H, Aliph), 3.95 (s, 3H, Aliph), 3.8 (m, 2H, Aliph), 3.30 (m, 4H, Aliph), 2.7 (m, 3H, Aliph), 2.07(m, 2H, Aliph), 2.1 (m, 1H, Aliph).

6h, Yield: 65%; mp: 164-167°; IR (KBr): 3433 (N-H stretch, amines), 3067 (C-H stretch, aromatic), 2926, 1442 (C-H stretch, Aliph), 1545 (C=N stretch), 1631 (C=O stretch, amide), 785 (C-Cl stretch), 694 (C-Br stretch); ¹H-NMR (CDCl₃): 8.5 (d, 1H, Ar), 8.15 (d, 2H, Ar), 7.45 (m, 4H, Ar), 7.26 (s, 1H, Ar), 7.20 (m, 2H, Ar), 5.32 (s, 1H, N-H, D₂O exchangeable), 4.1(s, 2H, Aliph), 3.8 (m, 2H, Aliph), 3.30 (m, 4H, Aliph), 2.7 (m, 3H, Aliph), 2.07(m, 2H, Aliph), 2.1 (m, 1H, Aliph).

Test compounds (6a-h) were evaluated in MCF7 breast, HOP62 lung, MIA-PA-CA-2 pancreatic, Colo205 and HT29 colon cancer cell lines (human) by following the sulforhodamine-B (SRB) semi automated assay protocol using adriamycin as a standard drug^[8]. The assay was carried out in a 96 well plate (each well of 100 µl capacity) using 90 µl of tumor suspension containing 5×10³ cells. Adriamycin and test compounds (6a-h) were tested at 15, 30, 60 and 120 µM concentrations. All experiments were done in triplicate. Results are average of three experiments. The average value was used to express activity as % inhibition as compared to control and then IC₅₀ value was determined. The results are given in Table 2.

Test compounds 6a-h, desloratadine and 10-bromodesloratadine were evaluated for FPT inhibition by following standard protocol^[1] and were tested at 30 and 60 µM concentrations. Desloratadine was used as a standard. Experiments were carried out in duplicate and the average value (% inhibition) was calculated relative to the vehicle control and then IC₅₀ value was determined. Data is given in Table 2.

TABLE 2: IC₅₀ VALUES FOR ANTICANCER EVALUATION AND FPT INHIBITION

Compound	IC ₅₀ Values in µM concentrations					
	Anticancer evaluation					FPT inhibition
	MCF	Hop62	Colo205	HT29	Miapaca-2	
6a	45	120	90	45	33	32
6b	22	60	90	<15	<15	34
6c	41	>120	105	45	45	<30
6d	>120	>120	>120	>120	>120	33
6e	27	53	<15	<15	37	40
6f	110	>120	>120	>120	>120	55
6g	70	>120	>120	65	60	60
6h	80	>120	>120	80	42	32
Desloratidine	25	50	85	20	25	60
10-Bromodesloratadine	<15	27	60	<15	18	53
Adriamycin	<15	<15	<15	<15	<15	--

Test compounds 6a-h in general exhibited dose dependent increase in anticancer activity in the cell lines but were less potent than adriamycin. Amongst the test compounds, 6e was most active with IC_{50} values of $<15 \mu\text{M}$ in two cell lines tested, while in remaining three cell lines its IC_{50} values were in the range of $27\text{-}53 \mu\text{M}$. Adriamycin had IC_{50} values of $<15 \mu\text{M}$ concentration. Adriamycin (doxorubicin) does not act by inhibiting FPT, but by interacting with DNA by intercalation^[9]. Hence it was not used as a standard in the FPT inhibition study.

Test compounds 6a-h exhibited potent FPT inhibitory activity at the doses studied and had IC_{50} values in the range of $32\text{-}40 \mu\text{M}$ except for 6f which had IC_{50} of $55 \mu\text{M}$. In comparison, desloratadine had IC_{50} of $60 \mu\text{M}$ and 10-bromodesloratadine had IC_{50} of $53 \mu\text{M}$. 10-bromodesloratadine was evaluated to examine the effect of introduction of bromine in desloratadine. The results show that 10-bromodesloratadine was more potent compared to desloratadine indicating that activity increases with bromination at C_{10} position.

It is clear that test compounds 6a-h in general are more potent than 10-bromodesloratadine and this enhanced activity can be attributed to the presence of zinc binding groups in these molecules.

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