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(54) Titre : UTILISATION DE COMPOSES NON HYDROPHOBES DANS LE TRAITEMENT DE METASTASES ET/OU DE
DEFAUTS DU CARTILAGE

(54) Title: NONE-HYDROPHOBIC COMPOUNDS FOR USE IN TREATING METASTASIS AND/OR CARTILAGE
DEFECT

(57) **Abrégé/Abstract:**

The present invention is directed to compounds, tautomers, stereoisomers, and chemically modified compounds thereof, and their use in preventing and/or treating tumors of metastasis and/or cartilage defect, and to a pharmaceutical composition comprising such compound.

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(54) Title: NONE-HYDROPHOBIC COMPOUNDS FOR USE IN TREATING METASTASIS AND/OR CARTILAGE DEFECT

(57) Abstract: The present invention is directed to compounds, tautomers, stereoisomers, and chemically modified compounds thereof, and their use in preventing and/or treating tumors of metastasis and/or cartilage defect, and to a pharmaceutical composition comprising such compound.

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None-hydrophobic compounds for use in treating metastasis and/or cartilage defect

5 The present invention refers to non-hydrophobic compounds, their use in prevention and/or treatment of metastasis and/or cartilage defect, and to a pharmaceutical composition comprising such compound, wherein the compound or composition is for example administered orally.

10 Technical background

Malignant melanoma is the skin cancer with the highest mortality rate. It is characterized by an early onset of metastasis formation and rapid disease progression; in the case of systemic metastases the 5-year survival rate is less than 10 % (Balch, C.M., et al., *Prognostic Factors Analysis of 17,600 Melanoma Patients: Validation of the*
15 *American Joint Committee on Cancer Melanoma Staging System*. Journal of Clinical Oncology, 2001. **19**(16): p. 3622-3634). In the Top5-EU-countries (Germany, UK, France, Spain, Italy), USA and Australia 125.000 incidences of primary tumors have been diagnosed in the year 2010; until 2025 an increase of 150.000 is expected (Globocan 2008). With the incidence of this malignancy increasing, new therapeutic attempts are
20 simultaneously emerging. They include the BRAF V600E inhibitor Vemurafenib, the c-Kit inhibitor Imatinib, the anti-CTLA-4 antibody Ipilimumab, which activates the immune system, as well as anti-PD1 immune checkpoint inhibitors Nivolumab and Lambrolizumab; however, overall survival remains poor (McArthur, G.A., et al., *Safety and efficacy of vemurafenib in BRAFV600E and BRAFV600K mutation-positive*
25 *melanoma (BRIM-3): extended follow-up of a phase 3, randomised, open-label study*. The Lancet Oncology, 2014. **15**(3): p. 323-332).

MIA, an 11 kDa protein secreted by melanocytic tumor cells, has been linked to cellular migration and invasion, which leads to the formation of metastases. It has also been
30 associated with immunosuppression of malignant melanoma (Jachimczak, P., et al., *Inhibition of immunosuppressive effects of melanoma-inhibiting activity (MIA) by antisense techniques*. International Journal of Cancer, 2005. **113**(1): p. 88-92) and is a diagnostic serum marker for melanoma disease progression as it is barely expressed in healthy tissue except by differentiating chondrocytes. MIA, also known as CD-RAP
35 (cartilage-derived retinoic acid-sensitive protein), is not only expressed in melanoma

cells but also in chondrocytes and modulates regeneration of regeneration of defect cartilage (Schmid R, Schiffner S, Opolka A, Grässel S, Schubert T, Moser M, Bosserhoff AK. Cell Death Dis. 2010 Nov 11). Analysis of the MIA/CD-RAP-*knockout*-mouse revealed that MIA/CD-RAP influences interactions between chondrocytes and their surrounding extracellular matrix, inhibits the proliferation of mesenchymal cells and promotes chondrocyte differentiation MIA is not only expressed in the context of melanoma but in numerous types of tumors, often late stage tumors such as breast cancer, glioma, pancreas carcinoma, and colon carcinoma etc. MIA allows the release of cells from the tumor by masking of binding sites for fibronectin and/or integrin and their migration in one direction which results in the invasion into other tissues and the formation of metastasis.

In WO 2011/113604 it is disclosed that MIA is functionally active as a homodimeric species and that peptides, e.g., AR71, are able to block the MIA-MIA interaction and in consequence to inhibit the formation of metastases and/or influence, cartilage formation, e.g., chondrocyte differentiation. No adverse effects were observed after a peptide treatment of mice. However, peptides generally make for poor drug candidates as they are readily degraded by proteases in the digestive tract or in the serum and are therefore typically not orally available.

Hence, the present invention is directed to improved inhibitors of the MIA dimerization for effective use in the prevention and/or treatment of metastasis and/or cartilage defect, which are for example even orally administrable.

Summary

In the present invention compounds have been investigated which are non-hydrophobic and interact with the dimerization site of MIA comprising or formed by for example at least three amino acid residues of the MIA protein selected from cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66, aspartate 67, leucine 76, tryptophan 102, aspartate 103, cysteine 106, valine 64, tyrosine 69, aspartate 87, lysine 91, glycine 54, leucine 58, phenylalanine 59, alanine 7, lysine 53, arginine 55, arginine 57, arginine 85 and lysine 94. Alternatively the dimerization site is selected from cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66, aspartate 67, leucine 76, tryptophan 102, aspartate 103, cysteine 106, alanine 7, lysine 53, arginine 55, arginine 57, arginine 85, and lysine 94. A further

selection is for example cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66, aspartate 67, leucine 76, tryptophan 102, aspartate 103 and cysteine 106.

The non-hydrophobic compounds are used alone or in combination for use in preventing
5 and/or treating metastasis caused by the dimerization of MIA protein and/or a cartilage defect, wherein regeneration is inhibited by MIA dimerization, wherein the compound(s) is/are selected from the group consisting of compound 1 to 270 of Table 1, tautomers, stereoisomers as well as chemically modified compounds thereof.

10 The metastasis preventable and/or treatable by the compounds of the present invention is based on any primary tumor expressing MIA such as melanoma, breast cancer, glioma, pancreas carcinoma, colon carcinoma, etc. The metastasis is located for example in the liver, lung, bone, colon, stomach, nerves, lymph nodes, skin and/or brain.

15 The compound of the present invention is administered alone or in combination with other non-hydrophobic compounds of the present invention, and in combination with a chemotherapeutic (e.g., Vermurafenib, Ipilimumab, Trametinib, Dabradenib, Dacarbazine, Paclitaxel, Carboplatin, Interferon-alpha, Aldesleukin etc.) comprising for example also an inhibitory cytokine (e.g., TGF alpha, TGF beta, interleukin etc.).

20

The compounds are either administered at the same time or consecutively.

The present invention is further directed to a pharmaceutical composition comprising one or more compounds of the present invention for use in preventing and/or treating of
25 metastasis caused by the dimerization of melanoma inhibitory activity (MIA) protein and/or a cartilage defect, wherein regeneration is inhibited by MIA dimerization, wherein the pharmaceutical composition comprises at least one compound selected from the group consisting of compound 1 to 270 of Table 1, tautomers, stereoisomers and chemically modified compounds thereof, and a pharmaceutically acceptable carrier
30 and/or solvent and optionally a chemotherapeutic.

In one embodiment the compound or the pharmaceutical composition of the present invention is administered orally.

Fig. 1 depicts a model of a MIA dimer wherein compounds of the present invention are bound to the MIA-MIA interaction site, also called MIA dimerization site.

Fig. 2 shows predicted passing of the blood-brain-barrier (**Fig. 2A**), predicted oral availability (**Fig. 2B**) and predicted systemic toxicity (**Fig. 2C**) of compound 3 (F1811-0131).

Fig. 3 presents inhibition of MIA dimerization by compounds 1 to 6 measured in the HTFP assay.

Fig. 4 shows Boyden chamber assays demonstrating the inhibition of melanoma cell migration by compounds 1, 2 and 3, respectively.

Fig. 5 depicts compounds 1 to 6 tested on proliferation and attachment of fibroblasts (**Fig. 5A, 5B**) and kidney cells HEK-293 (**Fig. 5C, 5D**), respectively. None of the tested compounds has an effect on these cells.

Fig. 6 presents the inhibition of the proliferation of the melanoma cell Mel-Im line by compound 1.

Detailed description

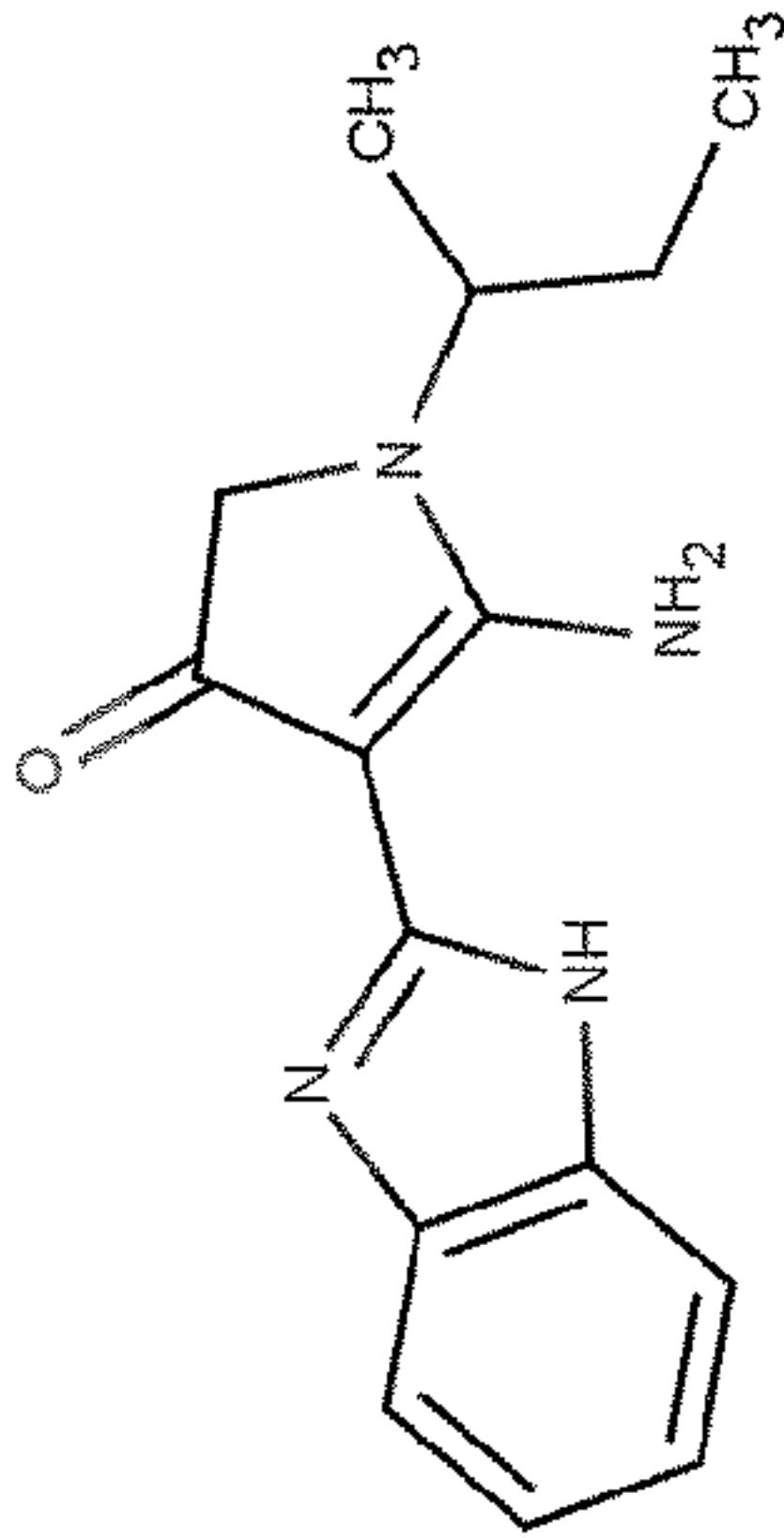
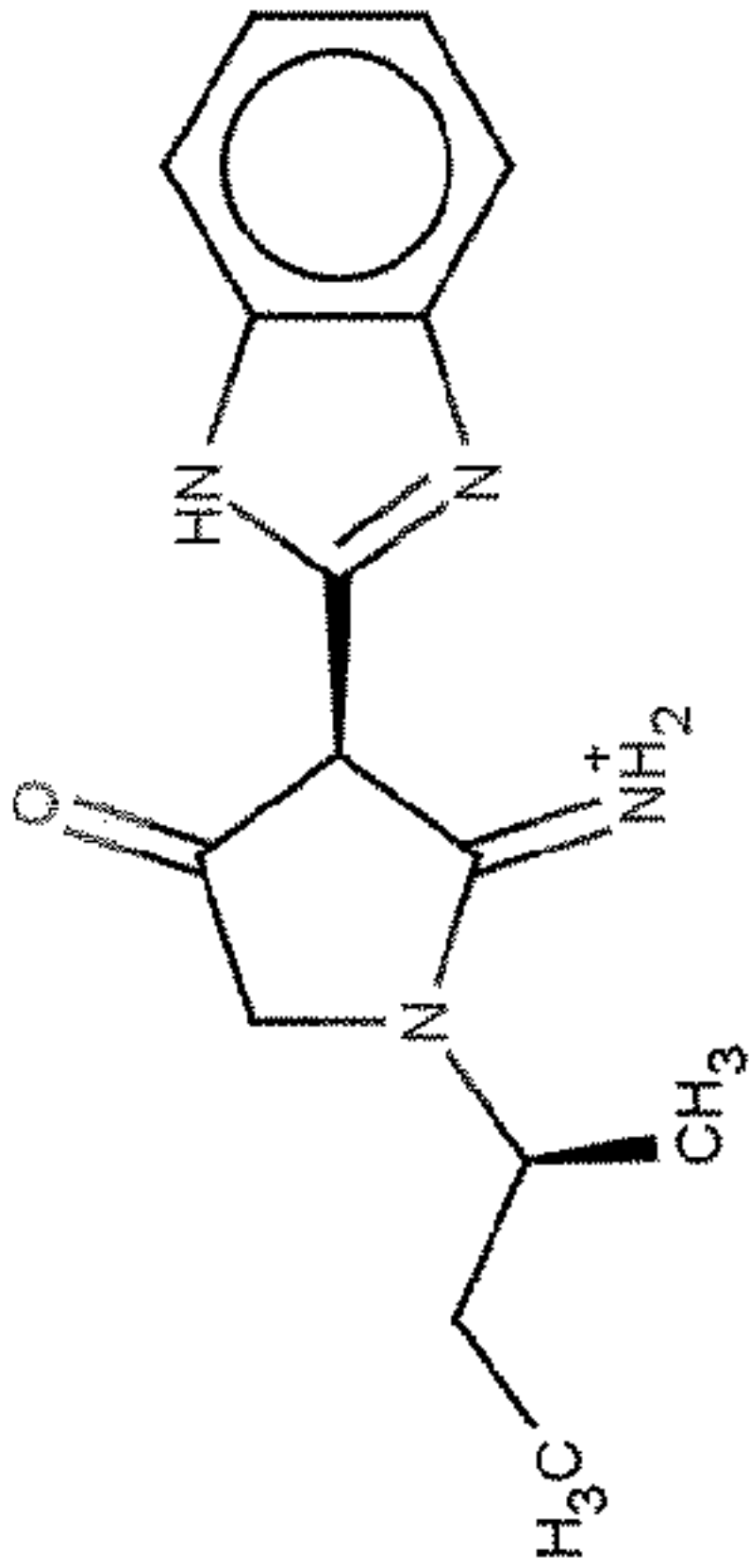
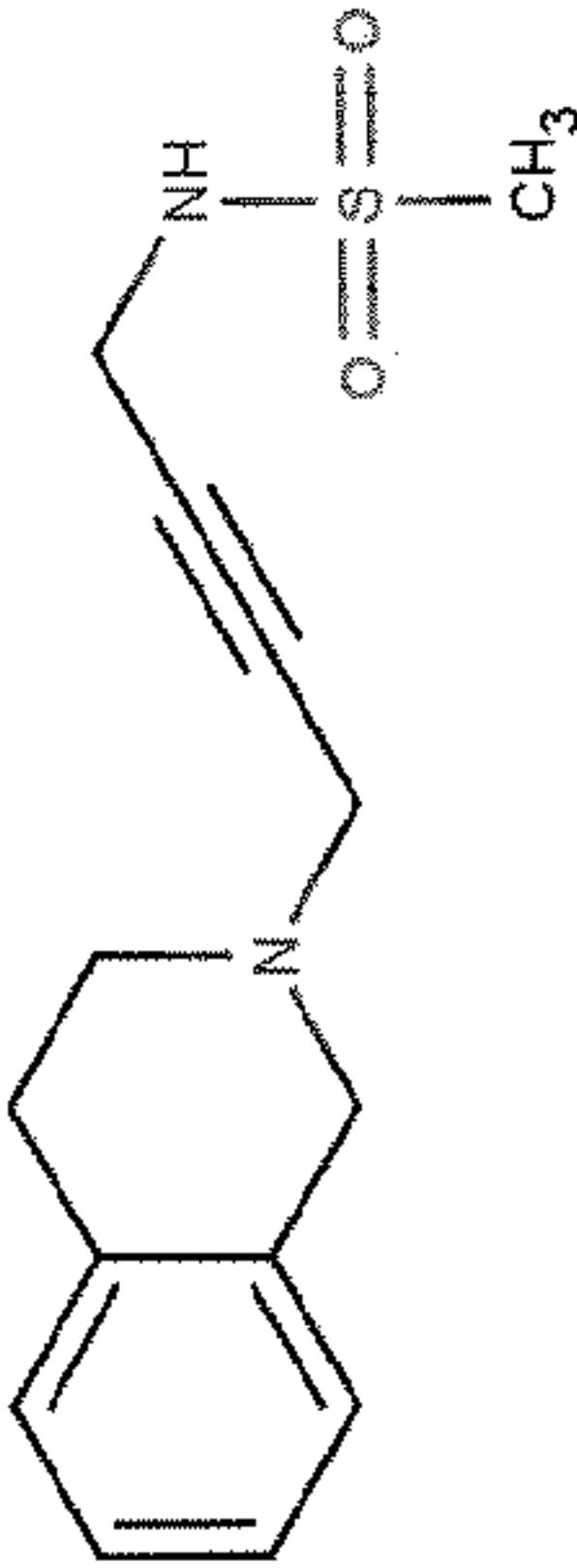
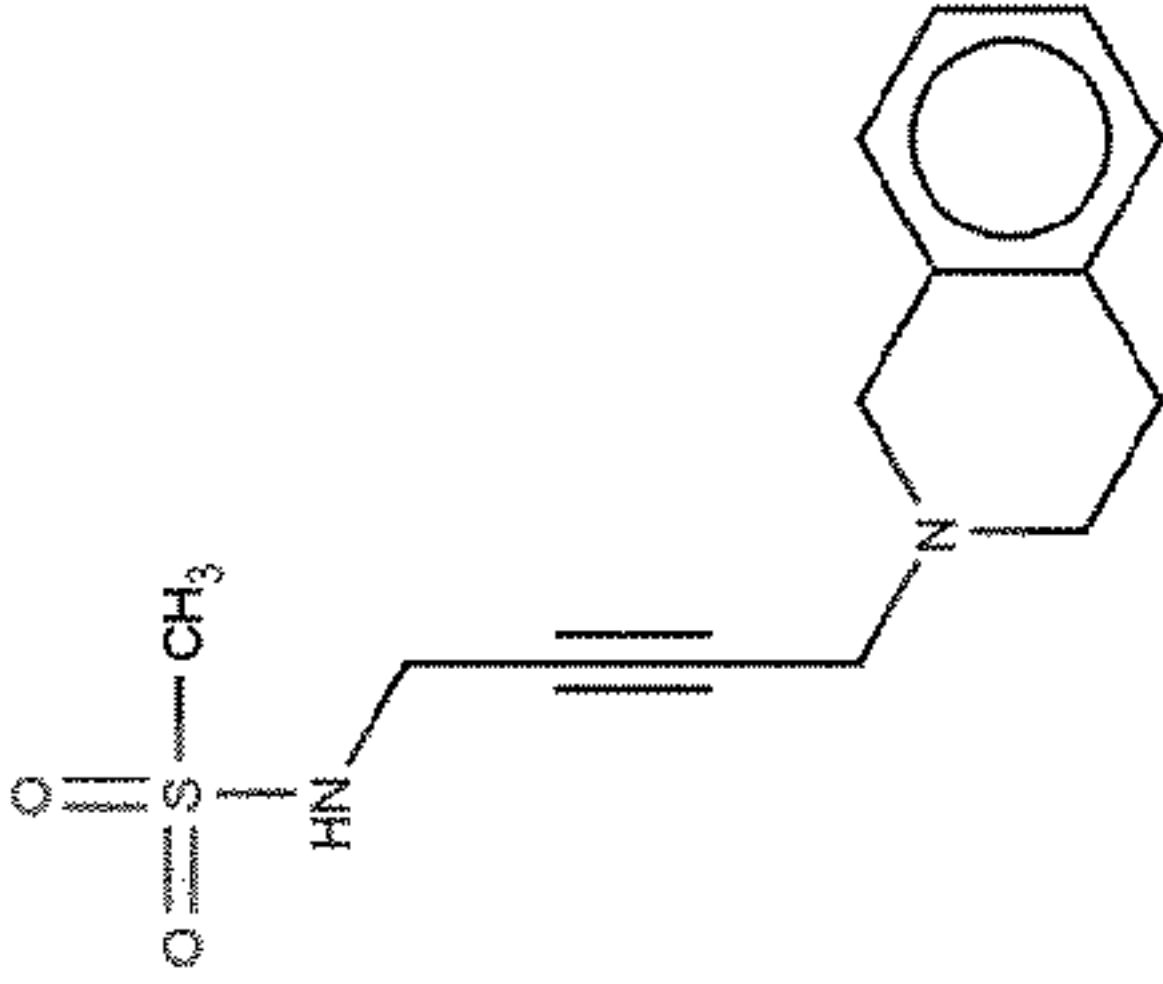
As a general treatment concept, the selective inhibition of MIA dimerization is an attractive therapy concept since, apart from its expression in malignant melanoma, MIA is only expressed by differentiating chondrocytes. Furthermore, the inhibition strategy of the present invention targets secreted extracellular MIA; thus the need for cell permeability of the compounds of the present invention is avoided. The compounds of the present invention are specifically developed and used for preventing and/or treating metastasis caused by the dimerization of melanoma inhibitory activity (MIA) protein or a cartilage defect which results for example in mechanical or immunological destruction and suppression of regeneration of the cartilage by MIA dimerization. MIA avoids the dedifferentiation of chondrocytes which would allow the proliferation of the chondrocytes. Without (dimerized) MIA, the chondrocytes transform into a certain type of precursor cells which are able to grow and to regenerate, i.e., close a cartilage defect. MIA blocks this regeneration.

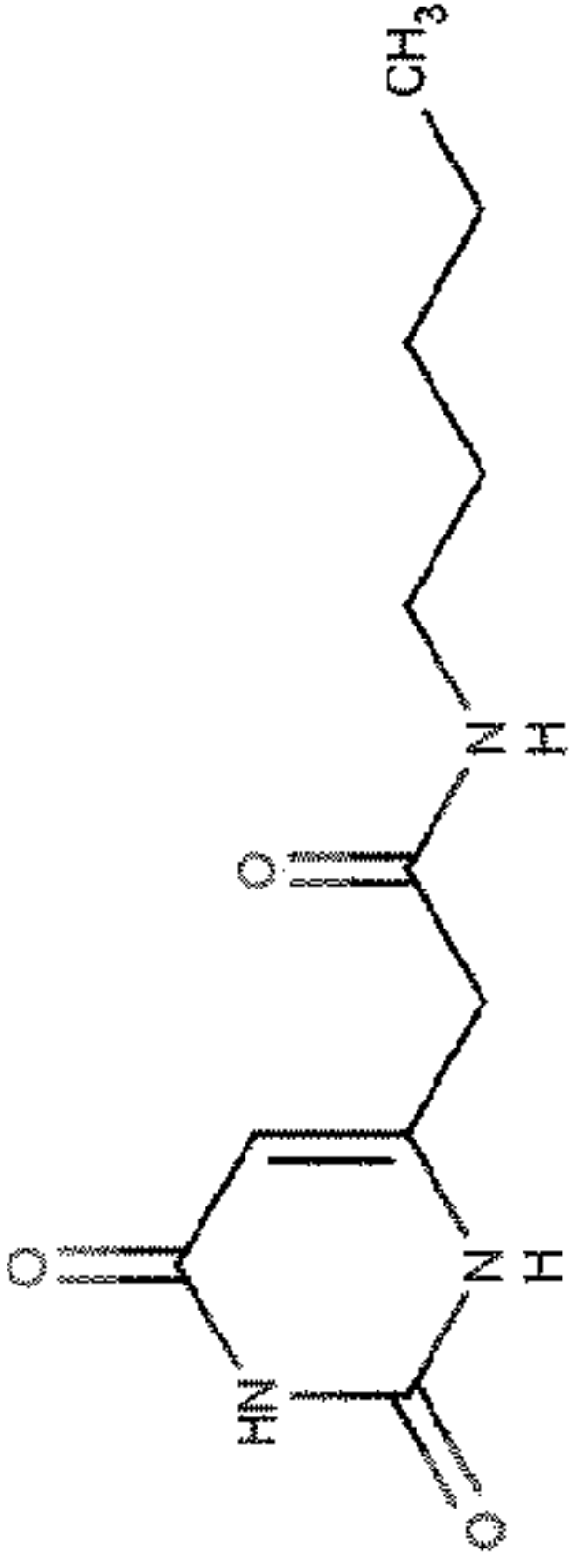
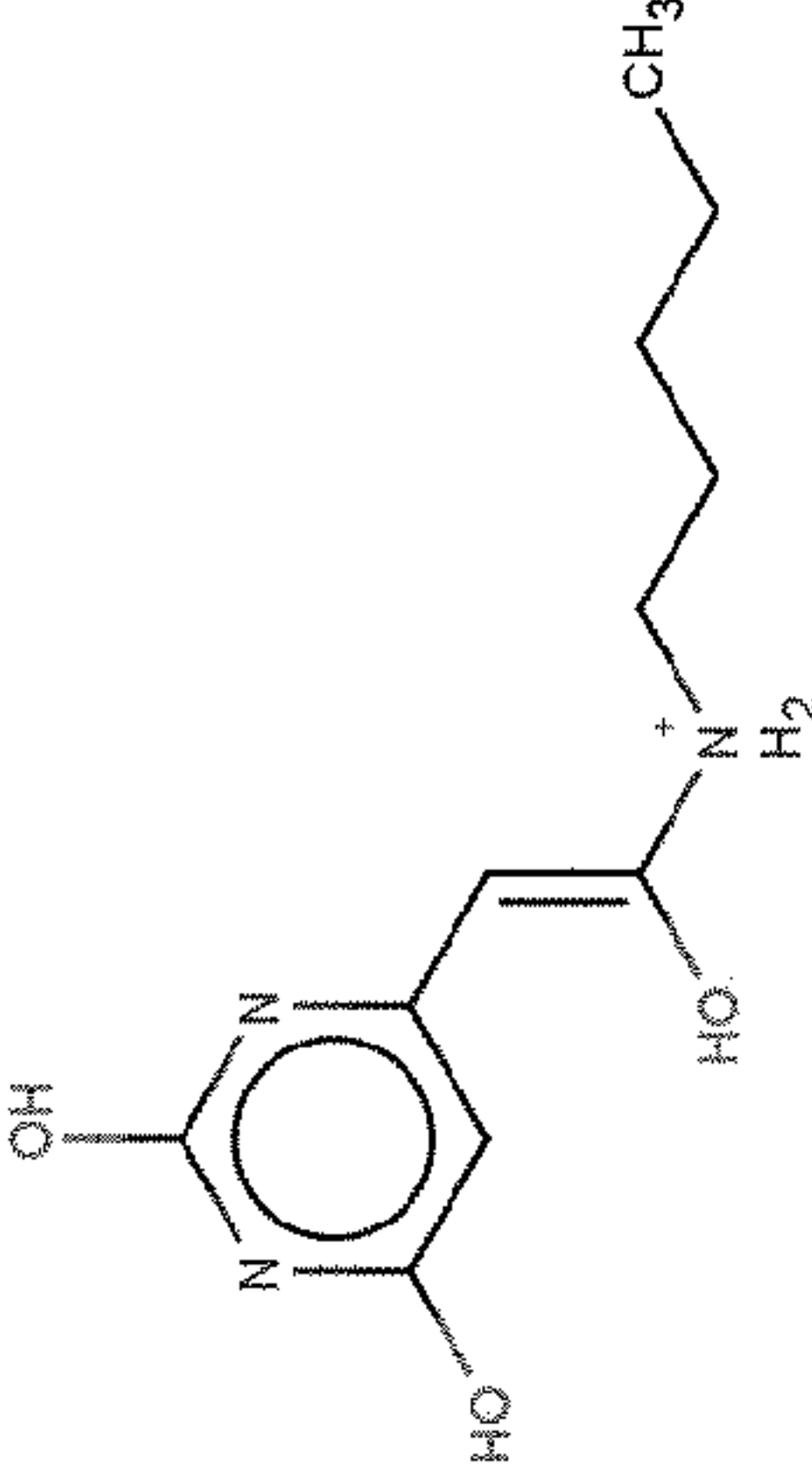
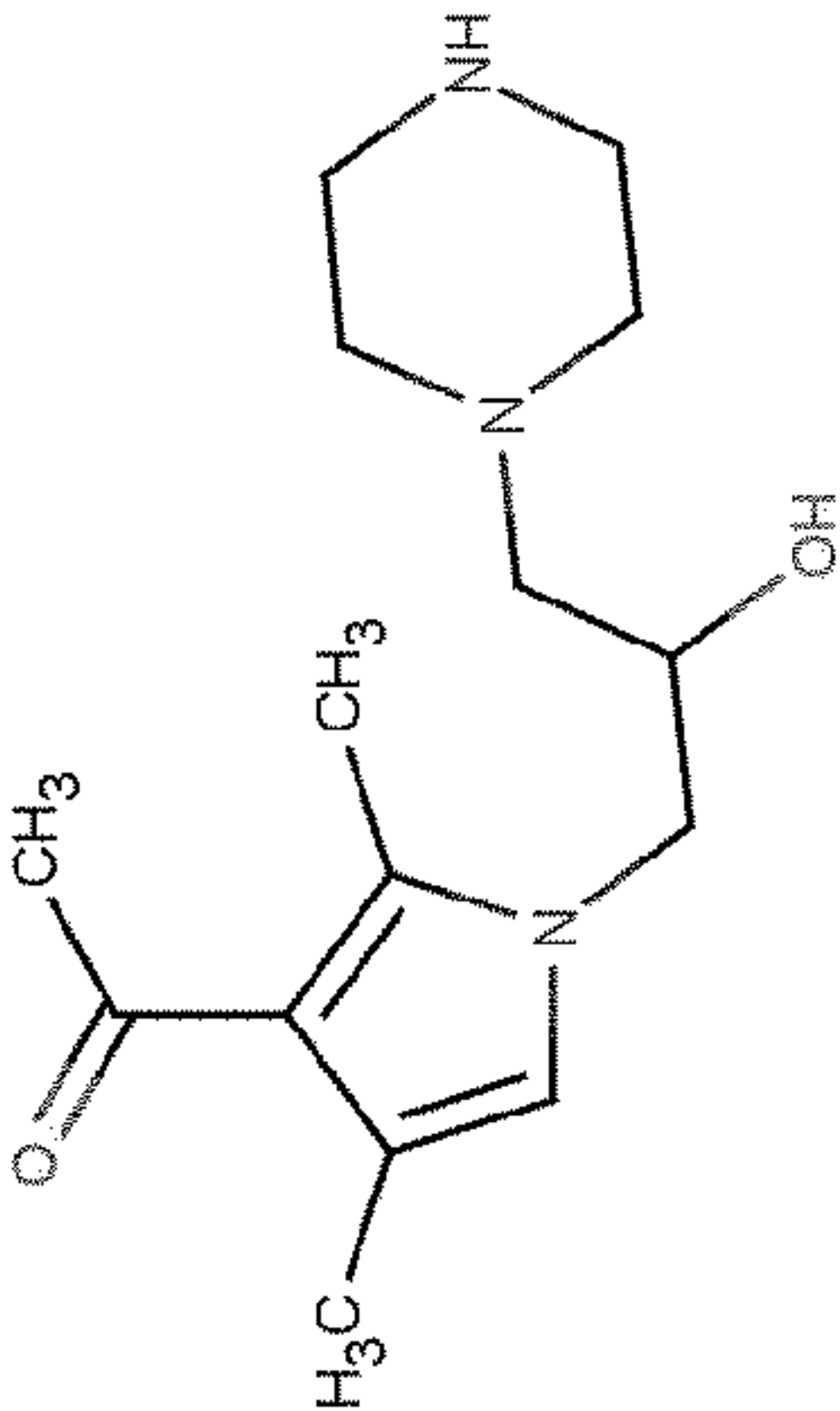
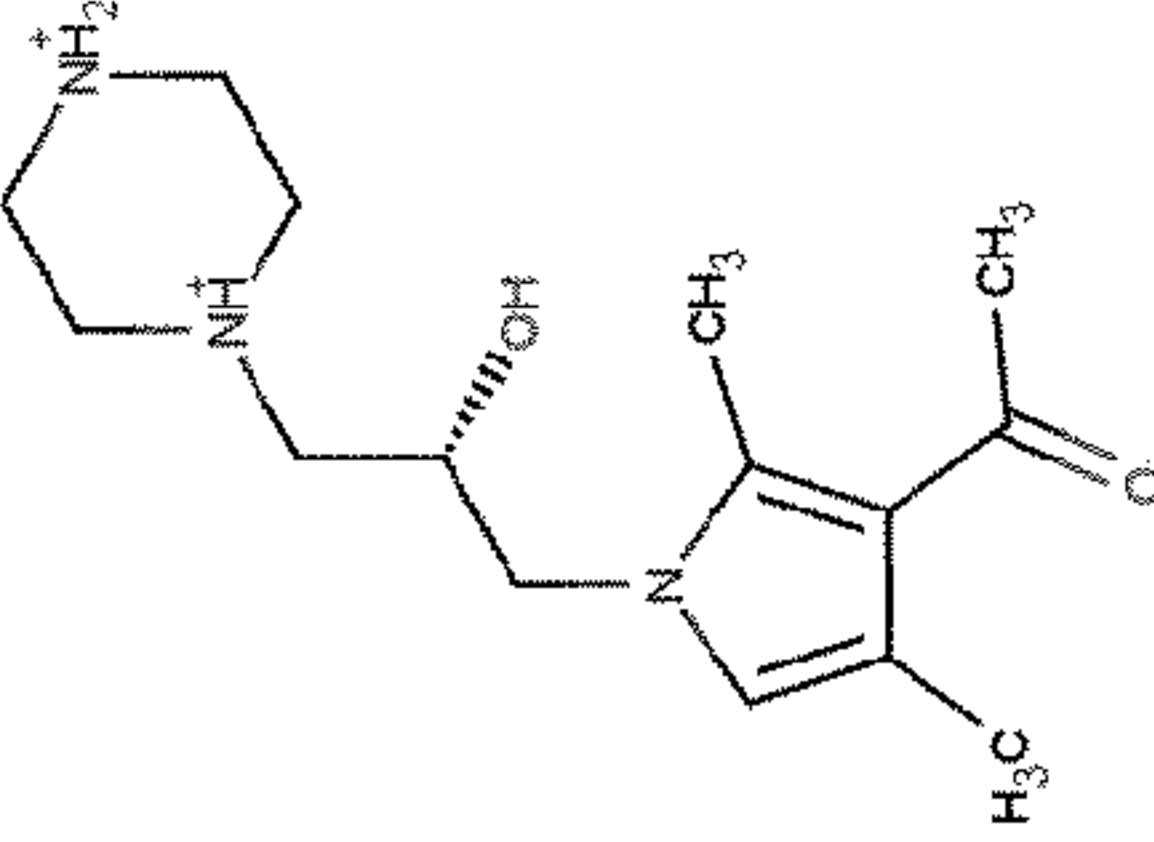
The compounds of the present invention interact and bind respectively, with/to the dimerization site of MIA proteins. Thus, the compounds of the present invention avoid the dimerization of MIA proteins and/or break up existing MIA dimers.

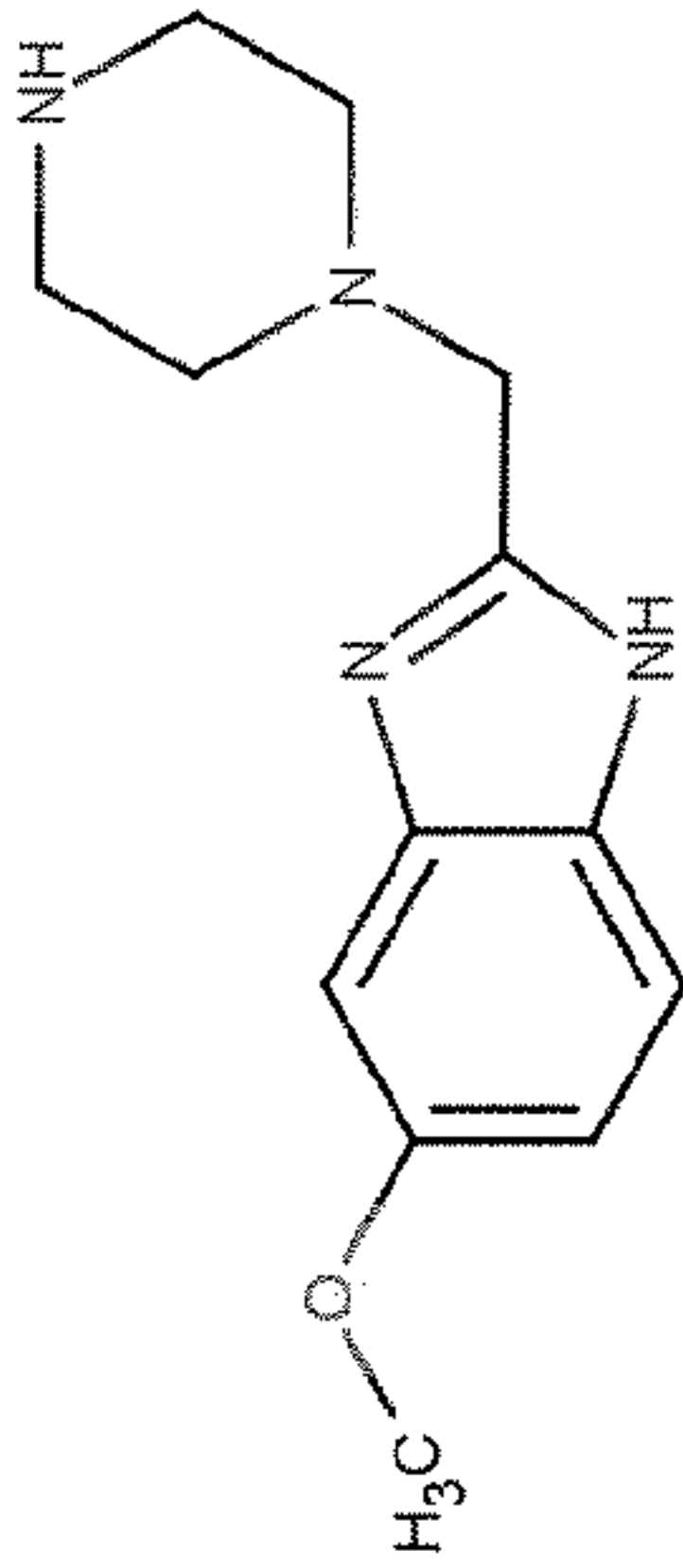
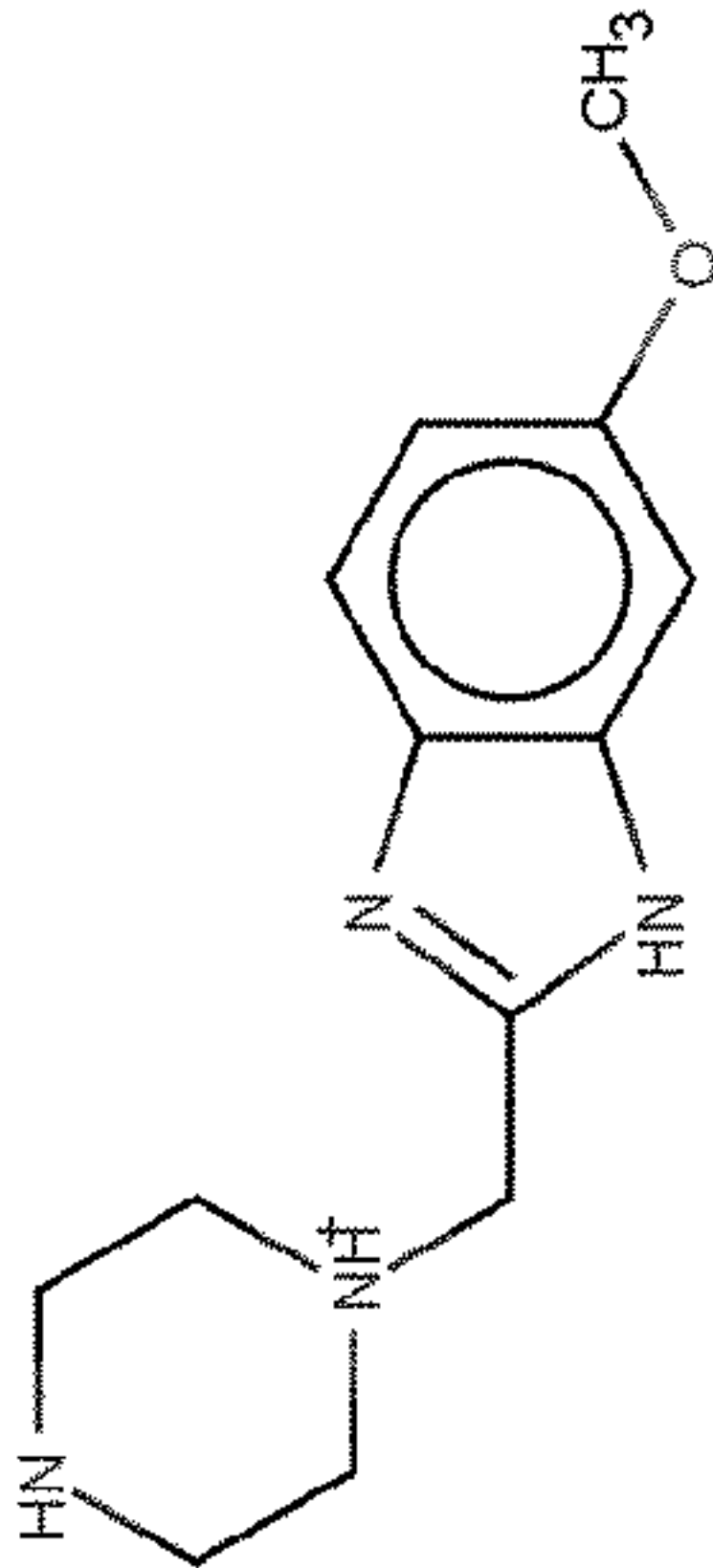
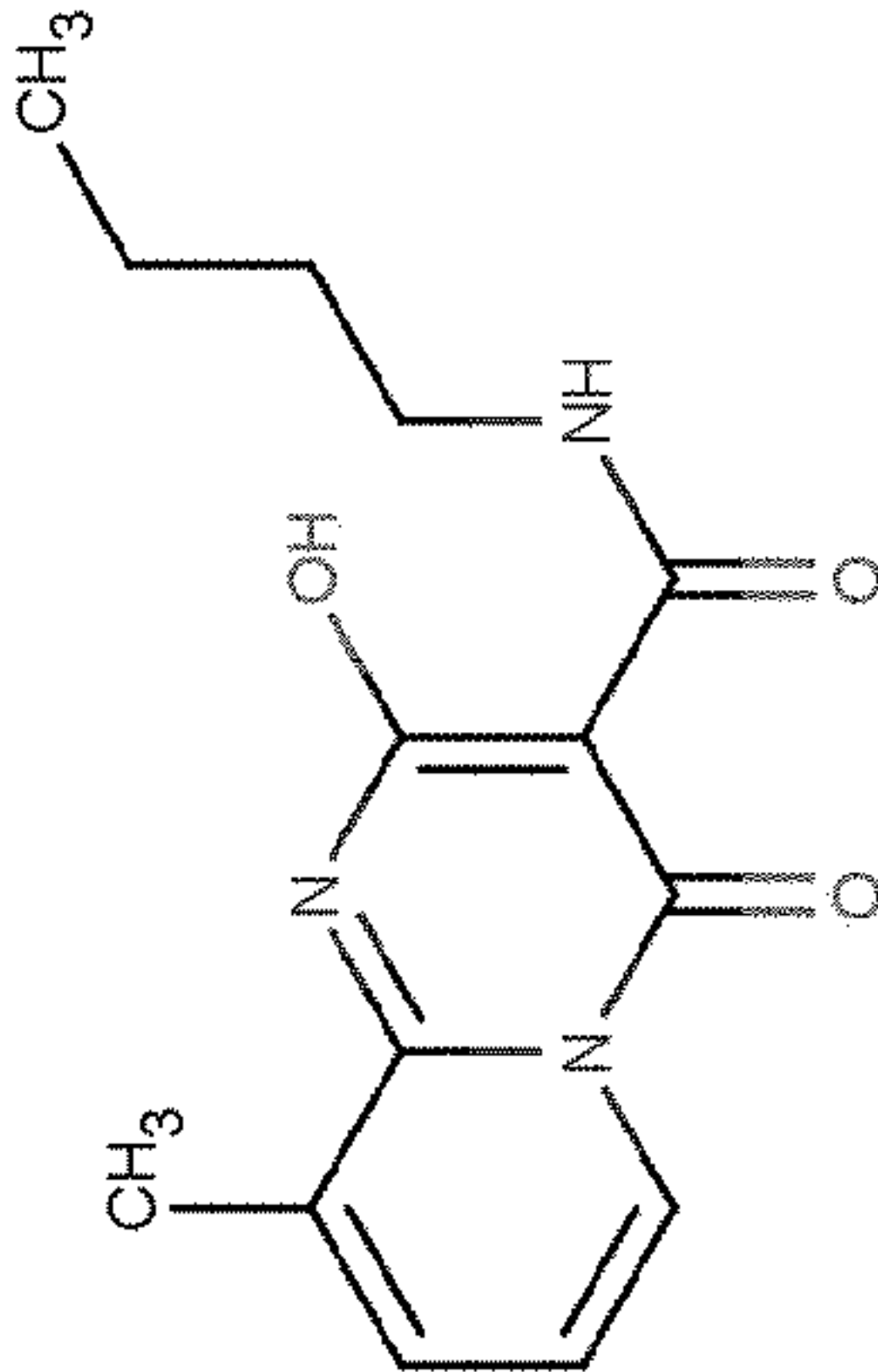
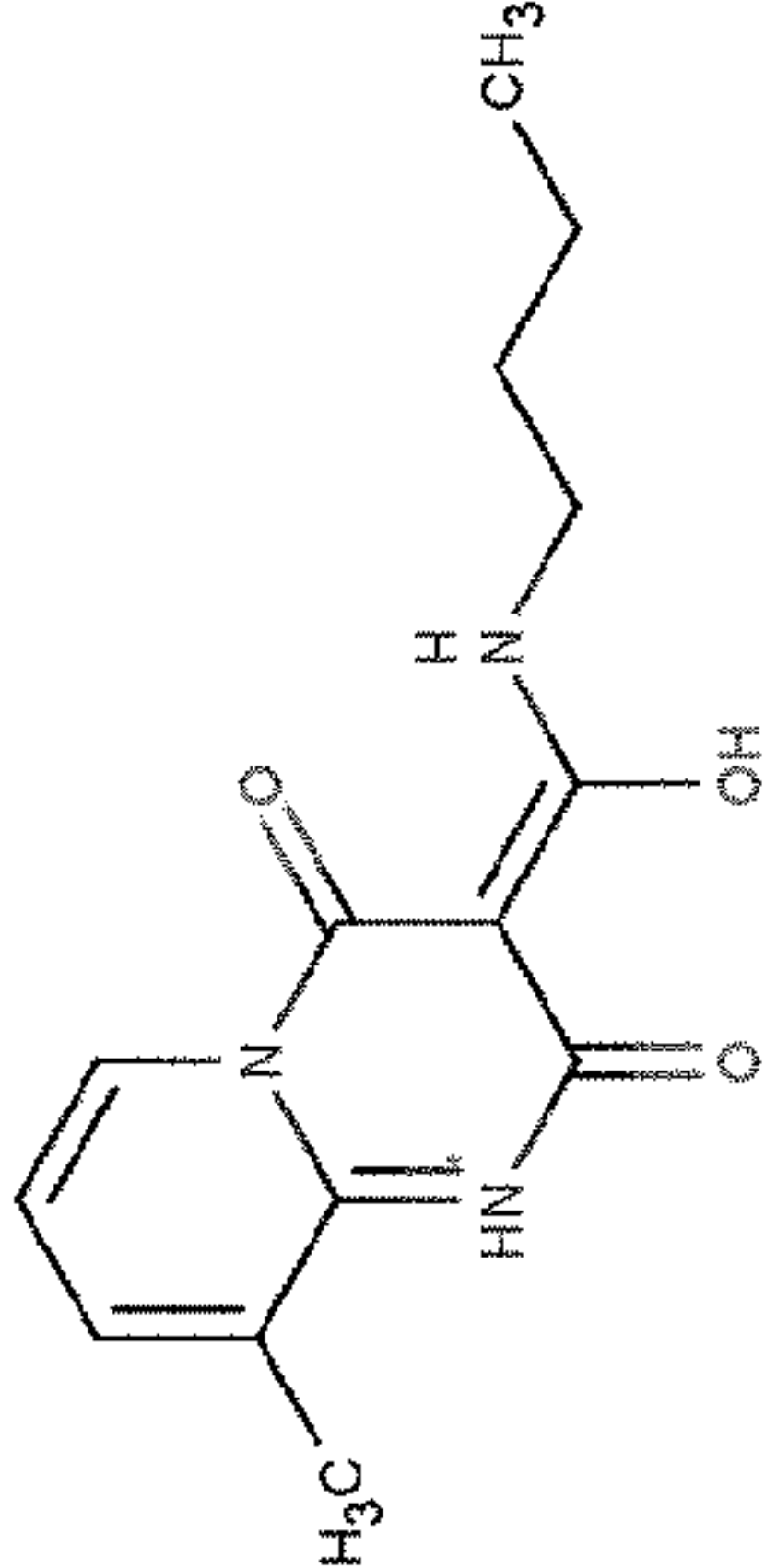
5 The MIA dimerization site which is the site where MIA proteins interact comprises or is formed by at least three amino acid residues of said MIA protein selected from cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66, aspartate 67, leucine 76, tryptophan 102, aspartate 103, cysteine 106, valine 64, tyrosine 69, aspartate 87, lysine 91, glycine 54, leucine 58, phenylalanine 59, alanine 7, lysine 53, arginine 55, arginine 57, arginine
10 85 and lysine 94. In one embodiment, the dimerization site of a MIA protein comprises or consists of cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66, aspartate 67, leucine 76, tryptophan 102, aspartate 103, cysteine 106, alanine 7, lysine 53, arginine 55, arginine 57, arginine 85, and lysine 94. In another embodiment, the MIA dimerization site comprises or consists of cysteine 17, serine 18, tyrosine 47, glycine 61, glycine 66,
15 aspartate 67, leucine 76, tryptophan 102, aspartate 103 and cysteine 106.

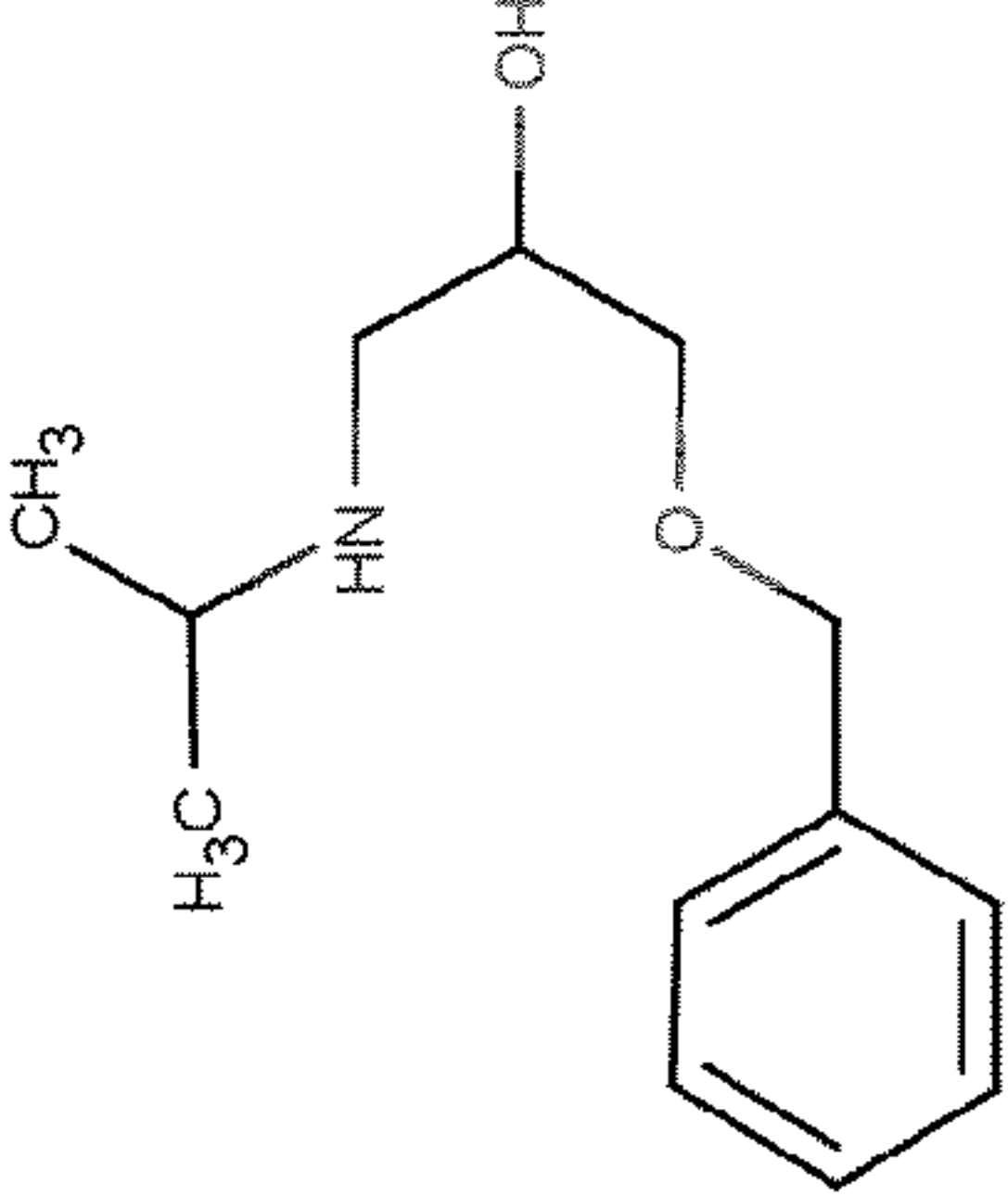
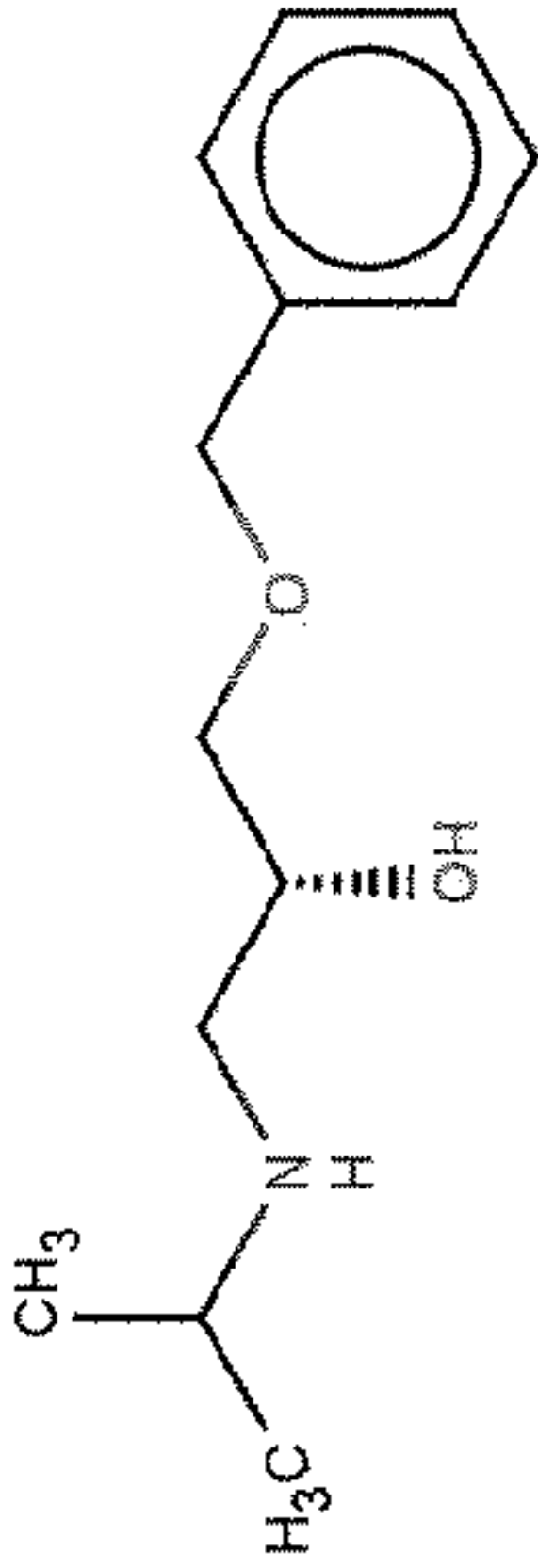
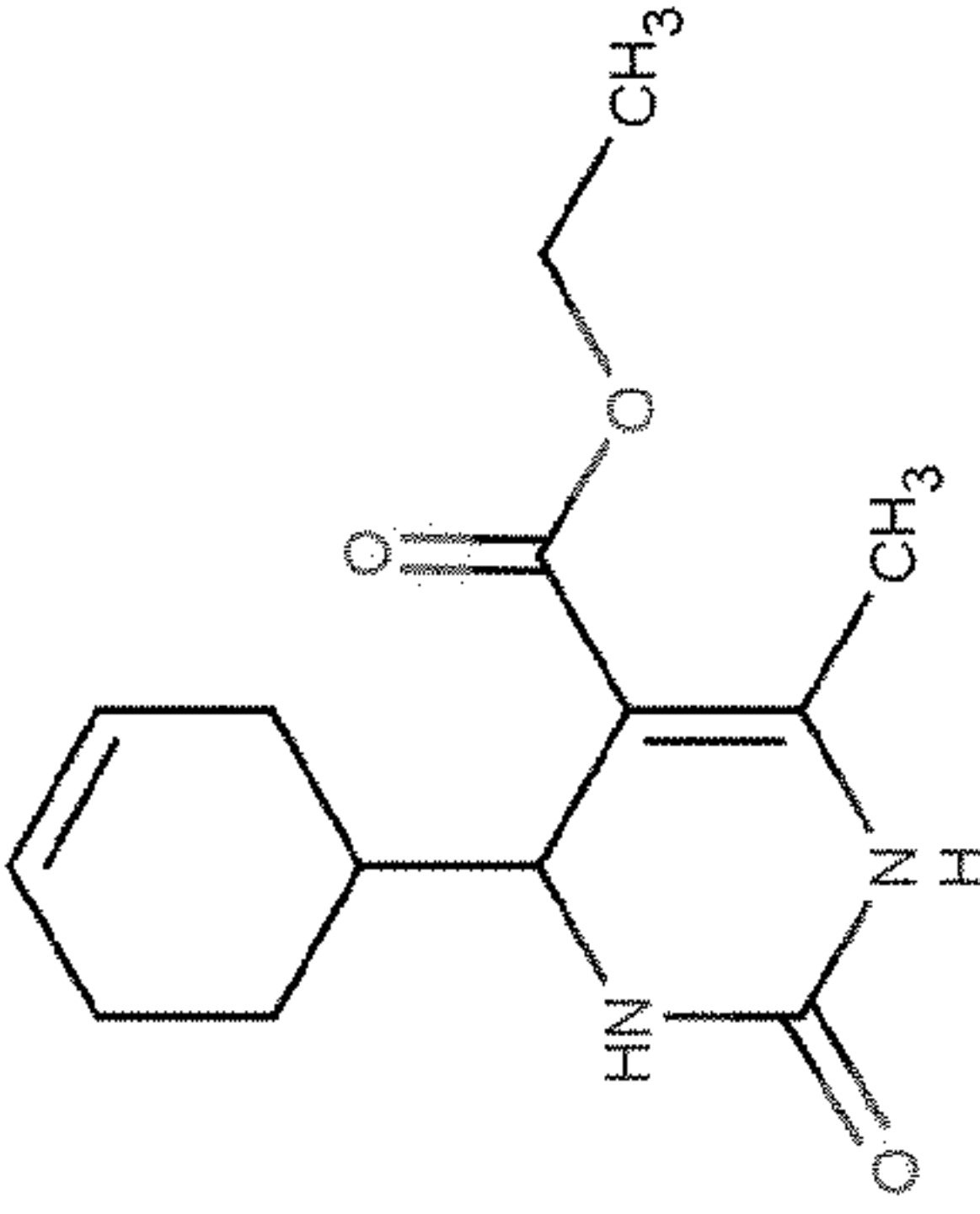
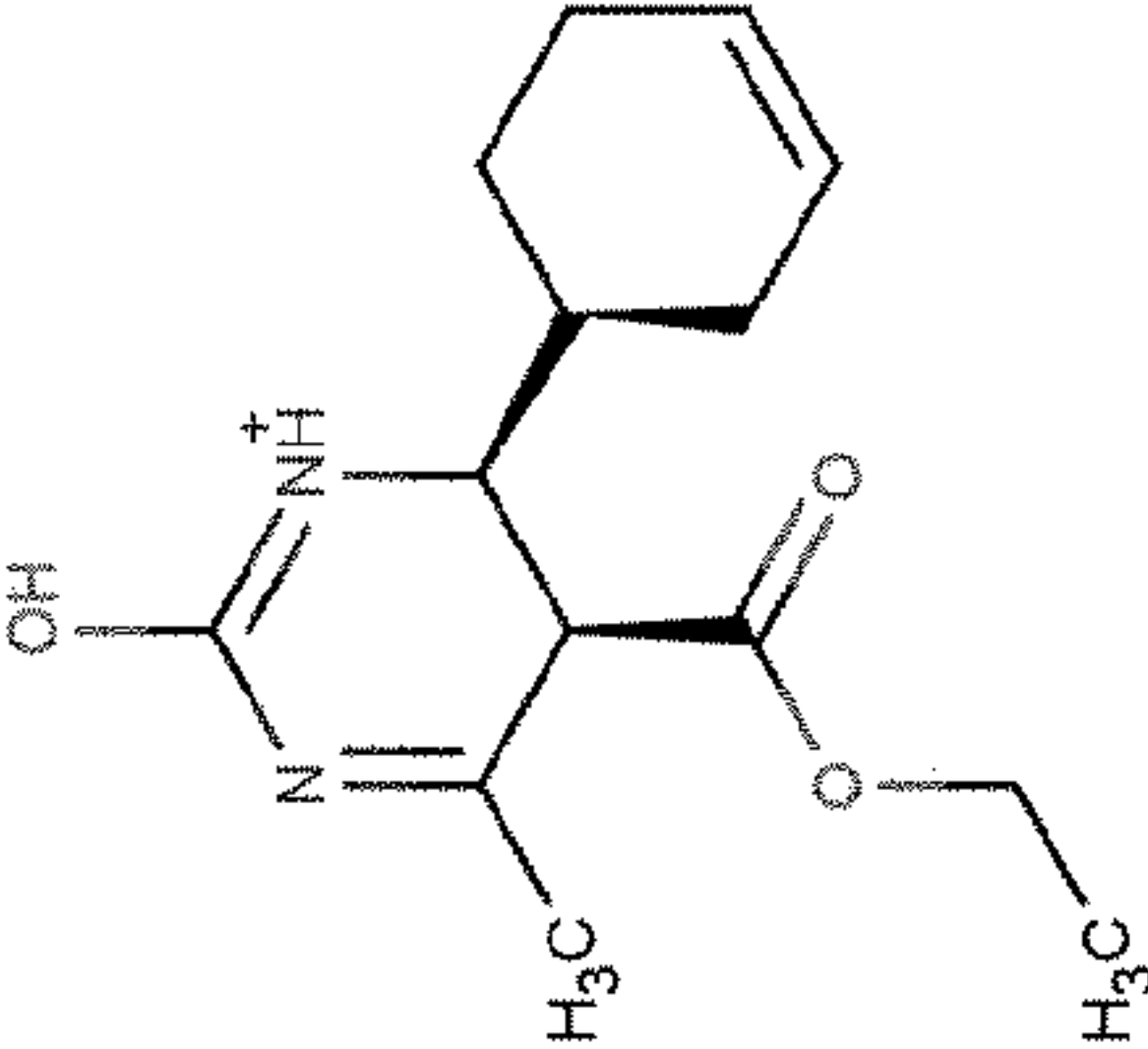
Via the dimerization site even more than two MIA proteins may interact and form an aggregate comprising or consisting of three or more MIA proteins. Fig. 1 presents a model of such MIA dimerization site virtually bound to numerous compounds of the
20 present invention. The amino acids of the MIA protein forming the dimerization site are hydrophobic in that the dimerization site is hydrophobic. Surprisingly, the compounds of the present invention interacting with the MIA dimerization site are non-hydrophobic. The compounds of the present invention for use in preventing and/or treating metastasis caused by the dimerization of MIA protein or a cartilage defect are listed in the following
25 Table 1. The invention further comprises the tautomers, stereoisomers and chemically modified compounds of the compounds listed in Table 1.

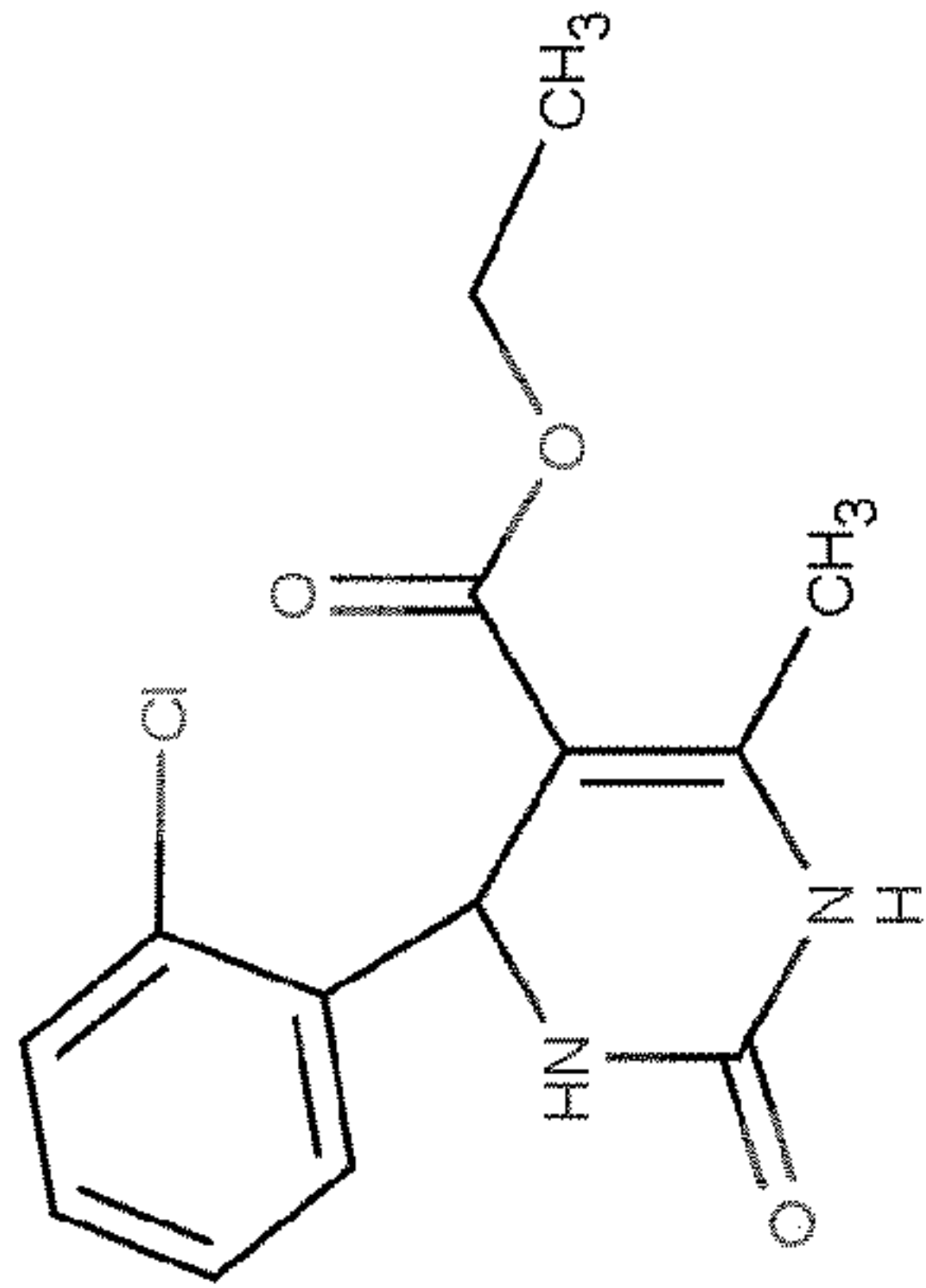
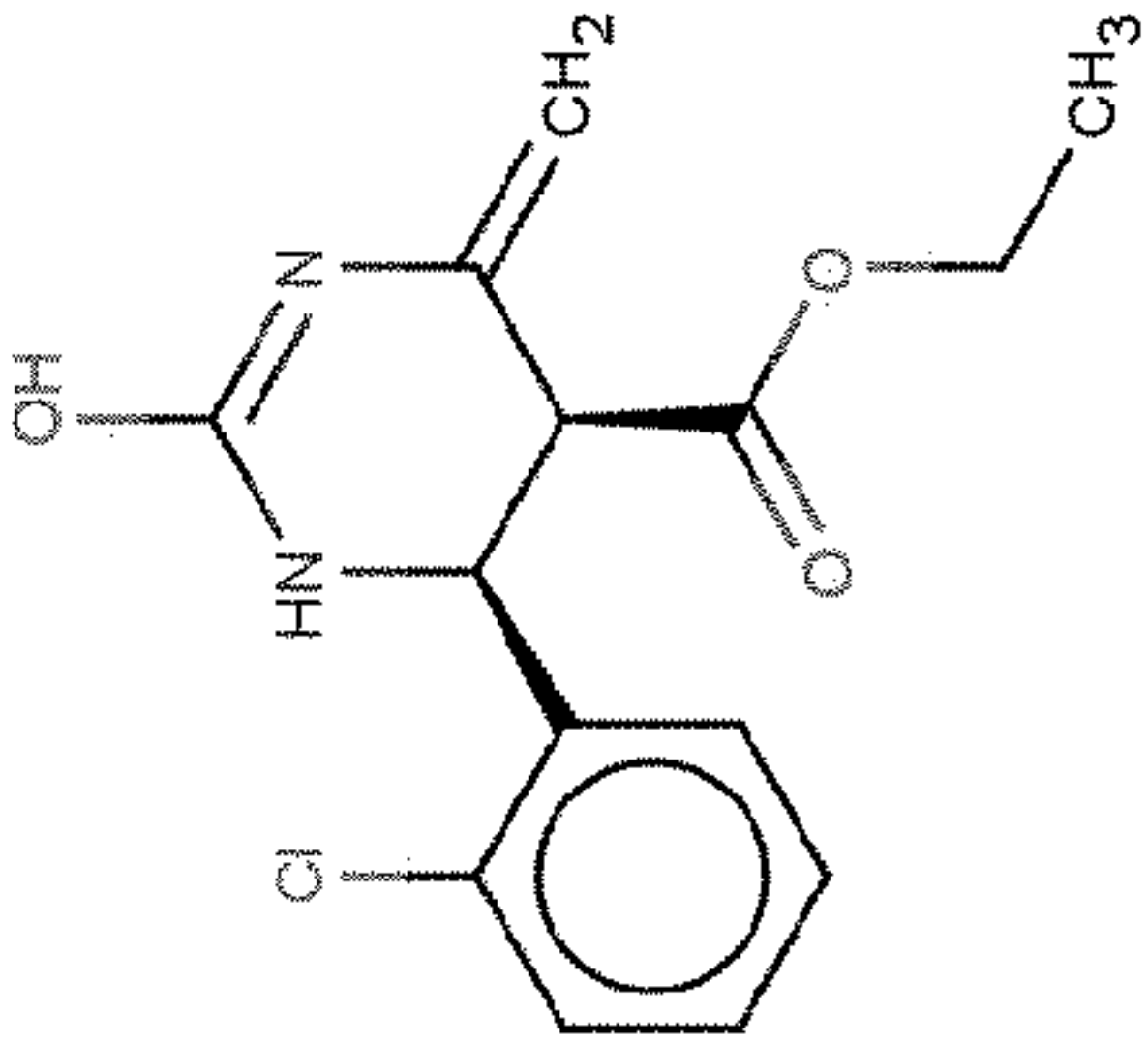
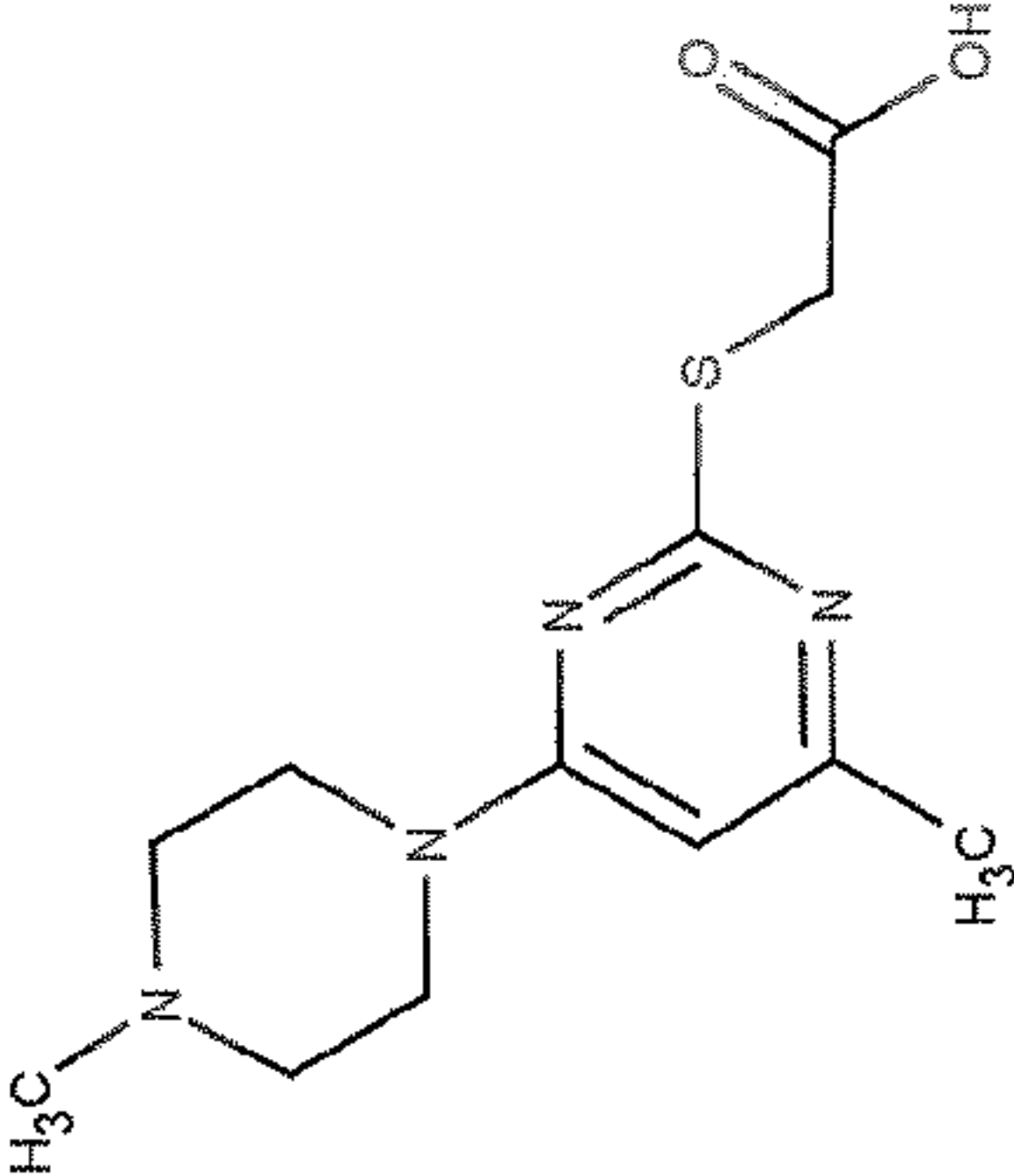
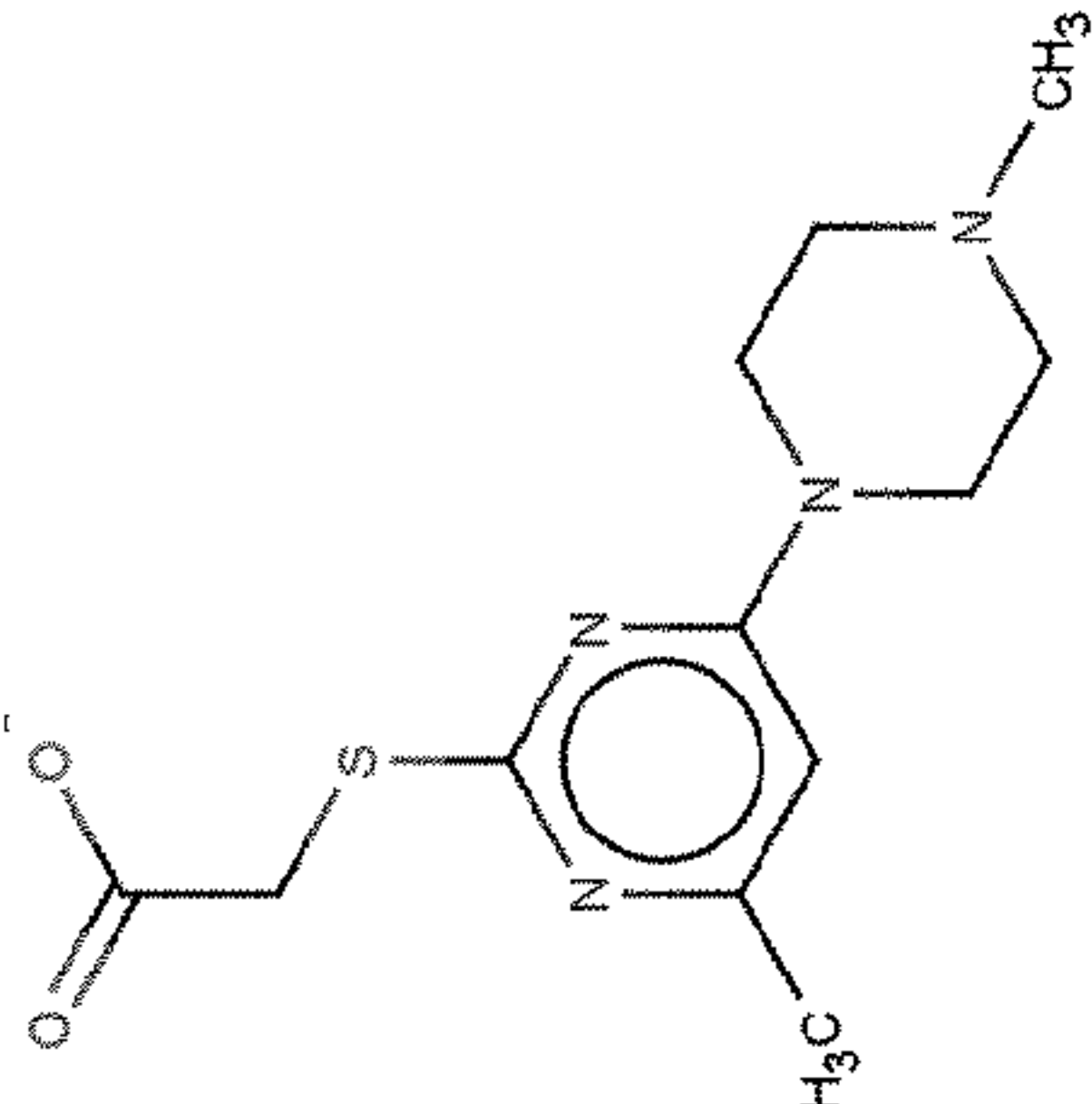
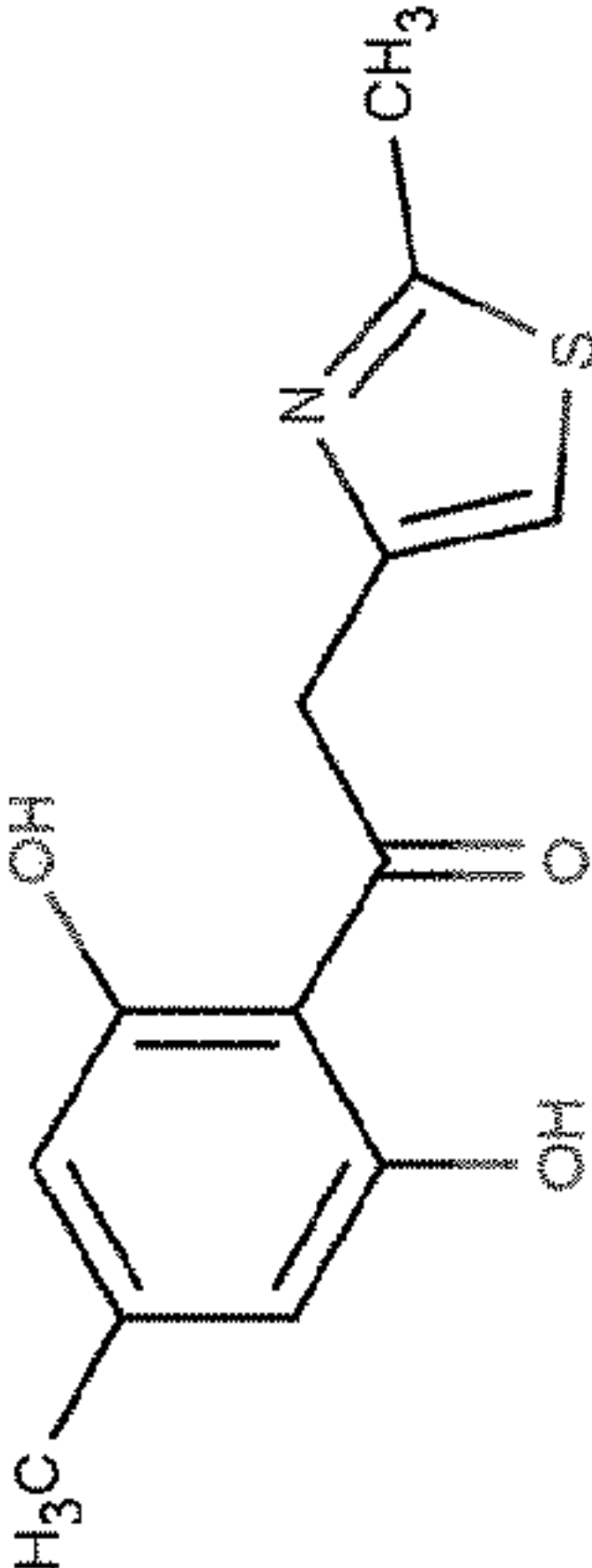
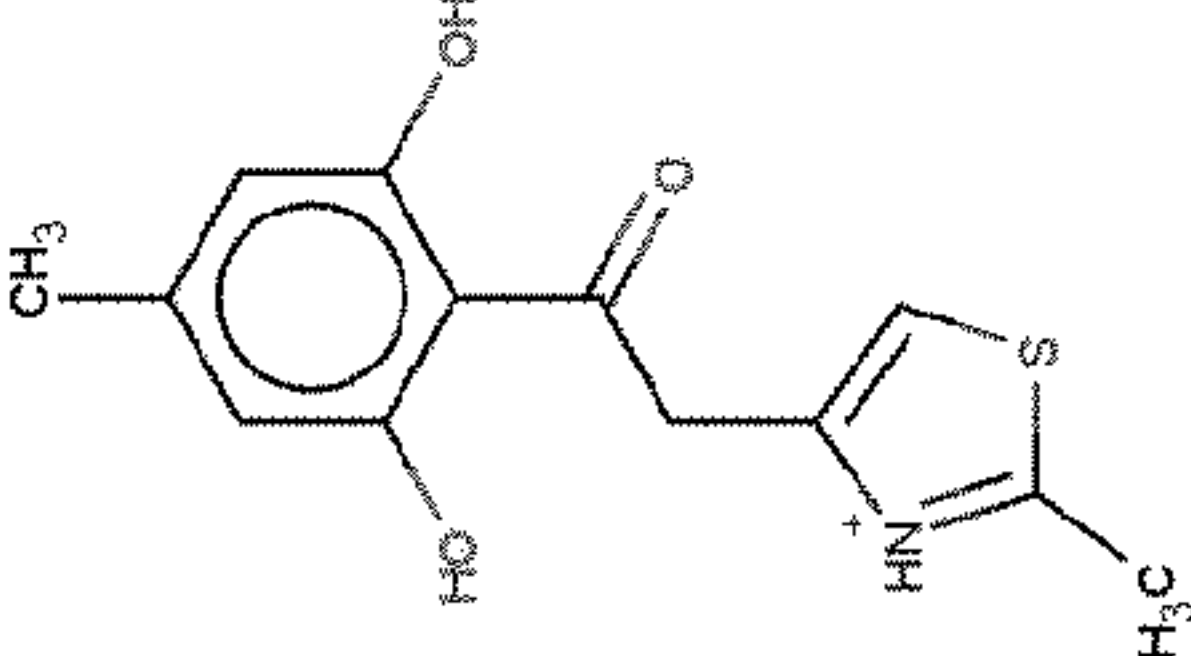
Table 1

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 1	 Preferred IUPAC Name = 5-amino-4-(1H-1,3-benzodiazol-2-yl)-1-(butan-2-yl)-2,3-dihydro-1H-pyrrol-3-one	F3225-8500		F3225-8500_8_2 (1 of 28)
Compound 2	 Preferred IUPAC Name = N-[4-(1,2,3,4-tetrahydroisoquinolin-2-yl)but-2-yn-1-yl]methanesulfonamide	F6244-3097		F6244-3097_1 (1 of 2)

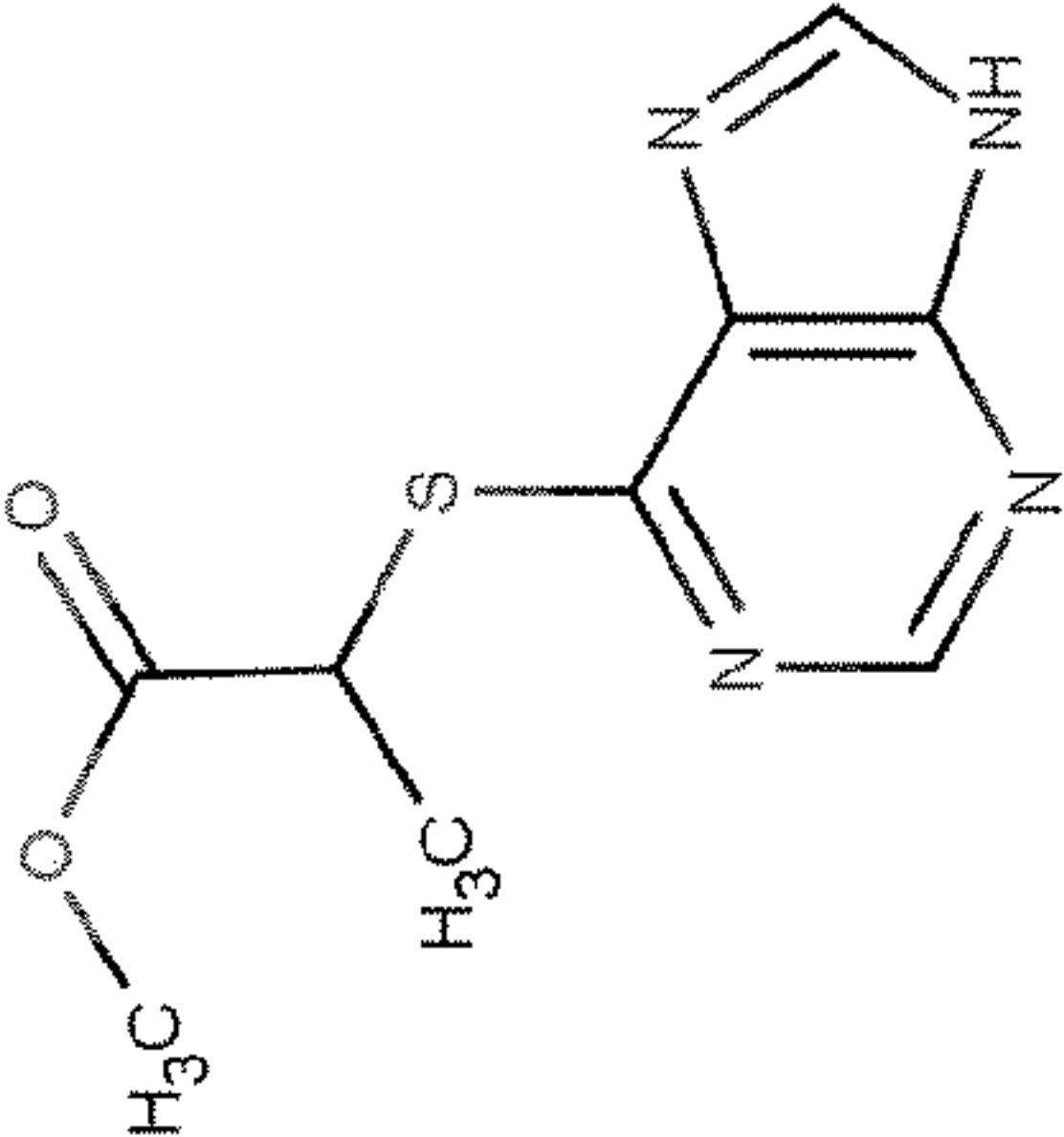
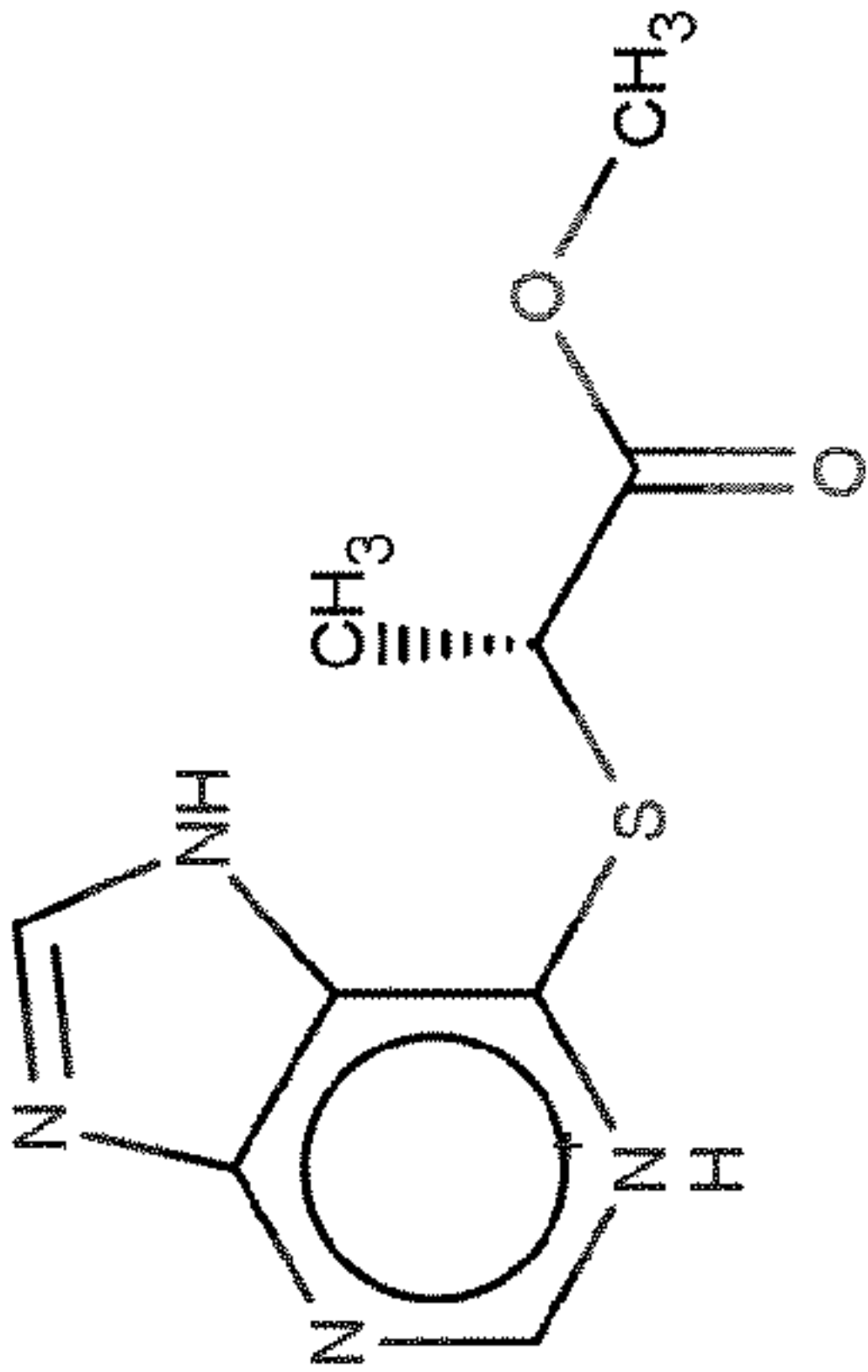
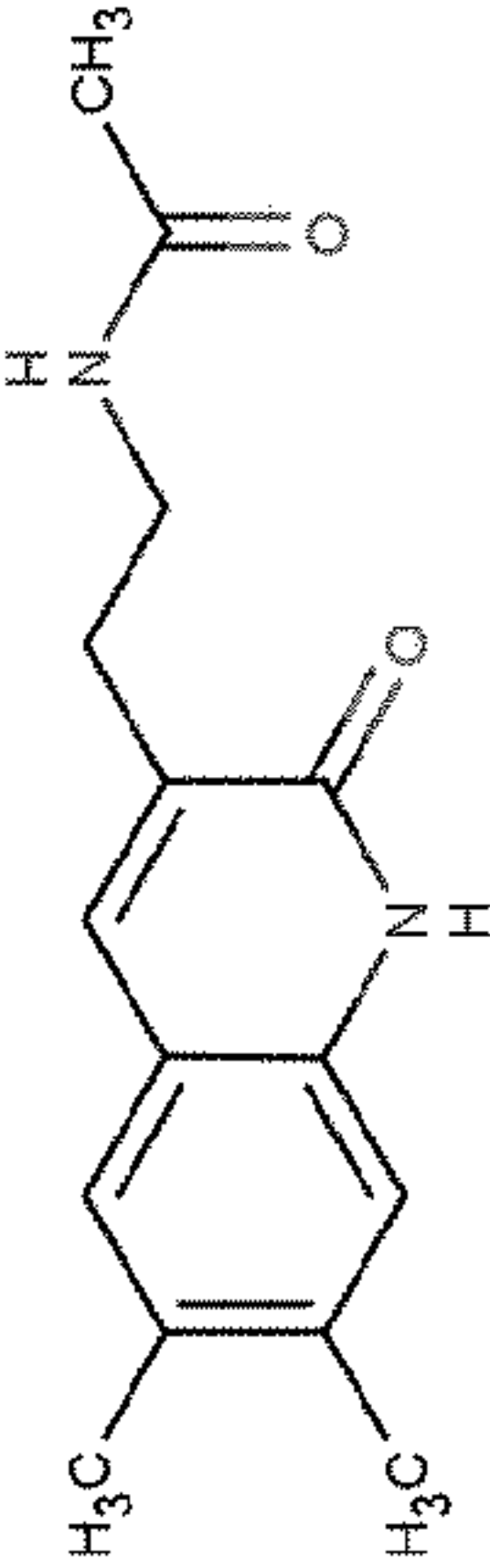
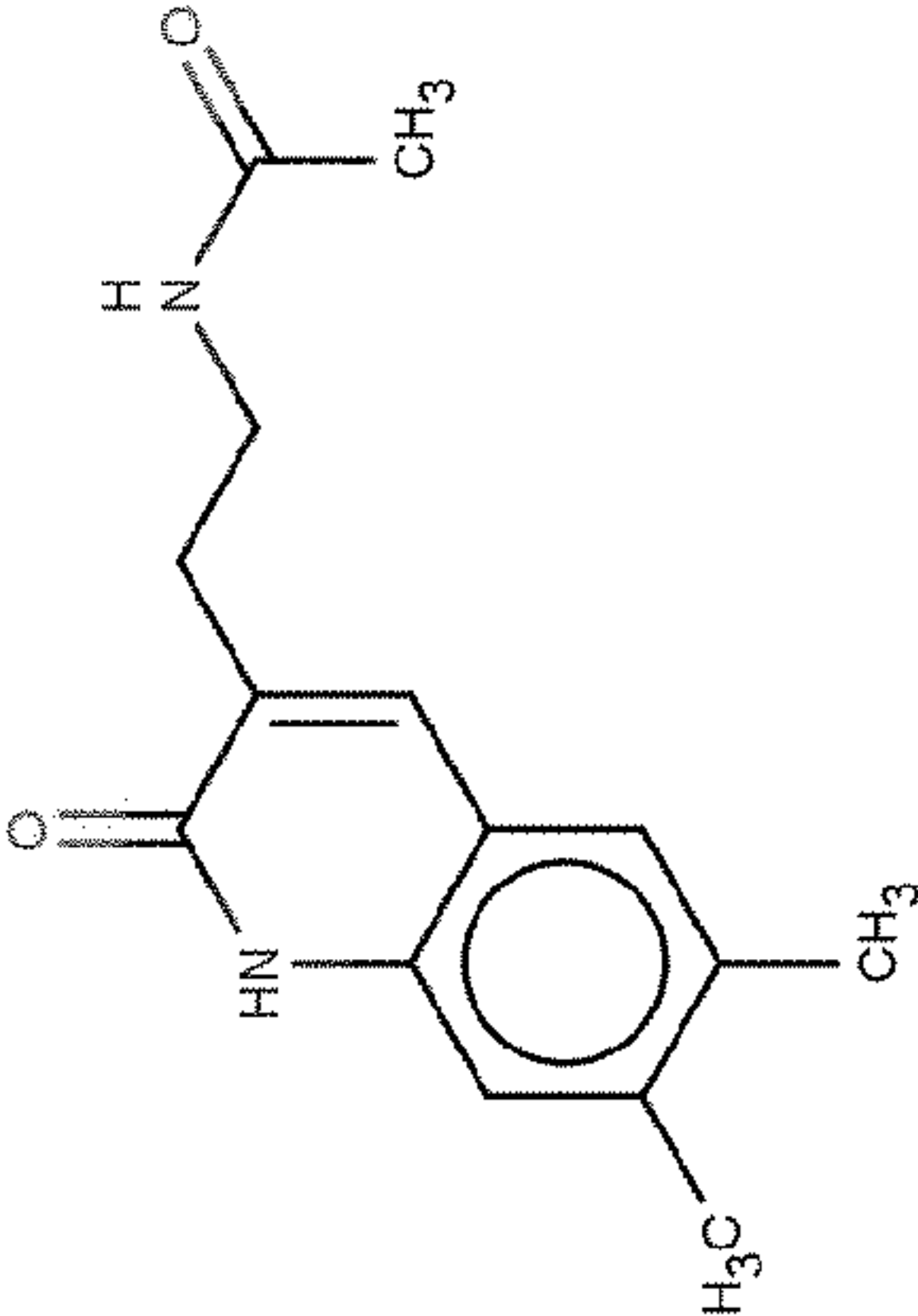
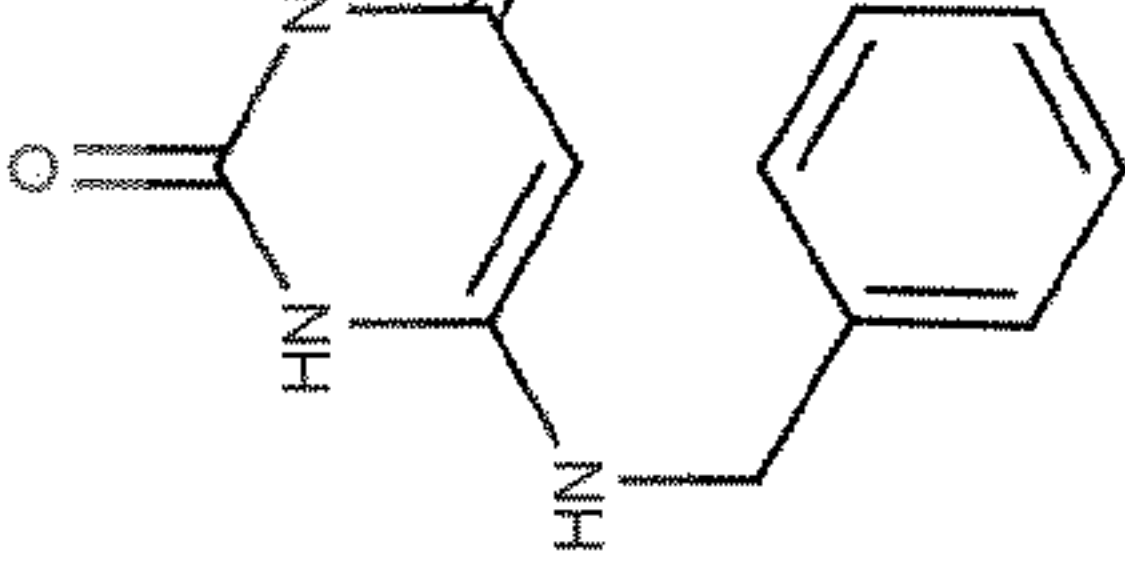
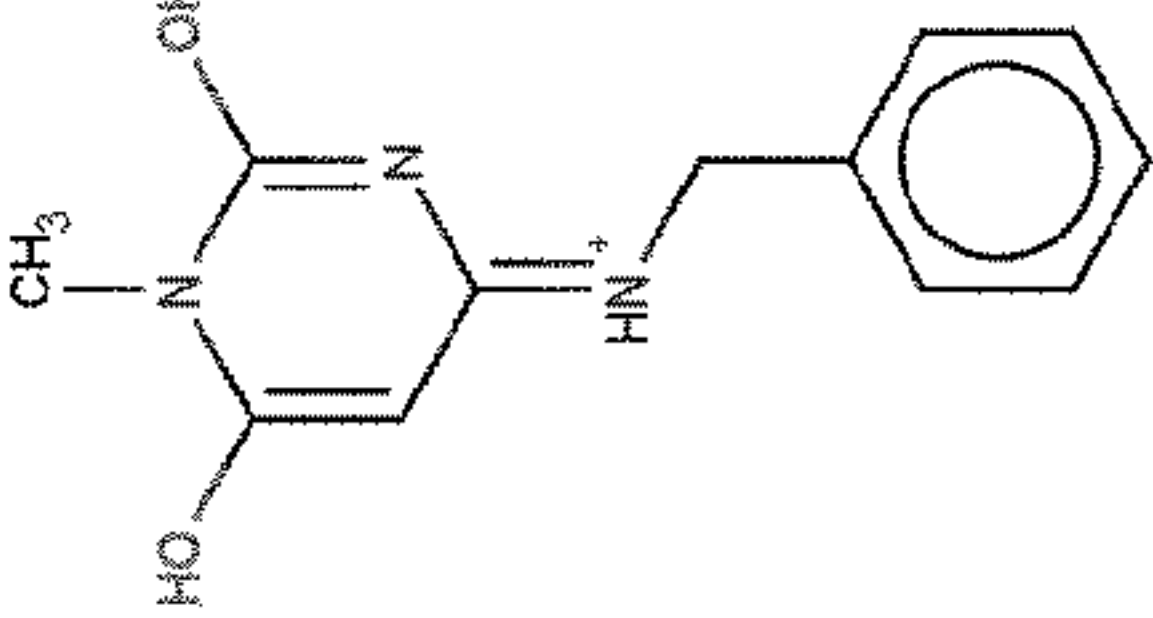
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 3	 Preferred IUPAC Name = 2-(2,6-dioxo-1,2,3,6-tetrahydropyrimidin-4-yl)-N-pentylacetamide	F1811-0131		F1811-0131_52_1 (1 of 76)
Compound 4	 Preferred IUPAC Name = 1-{1-[2-hydroxy-3-(piperazin-1-yl)propyl]-2,4-dimethyl-1H-pyrrol-3-yl}ethan-1-ol	F3255-0159		F3255-0159_3_2 (1 of 8)

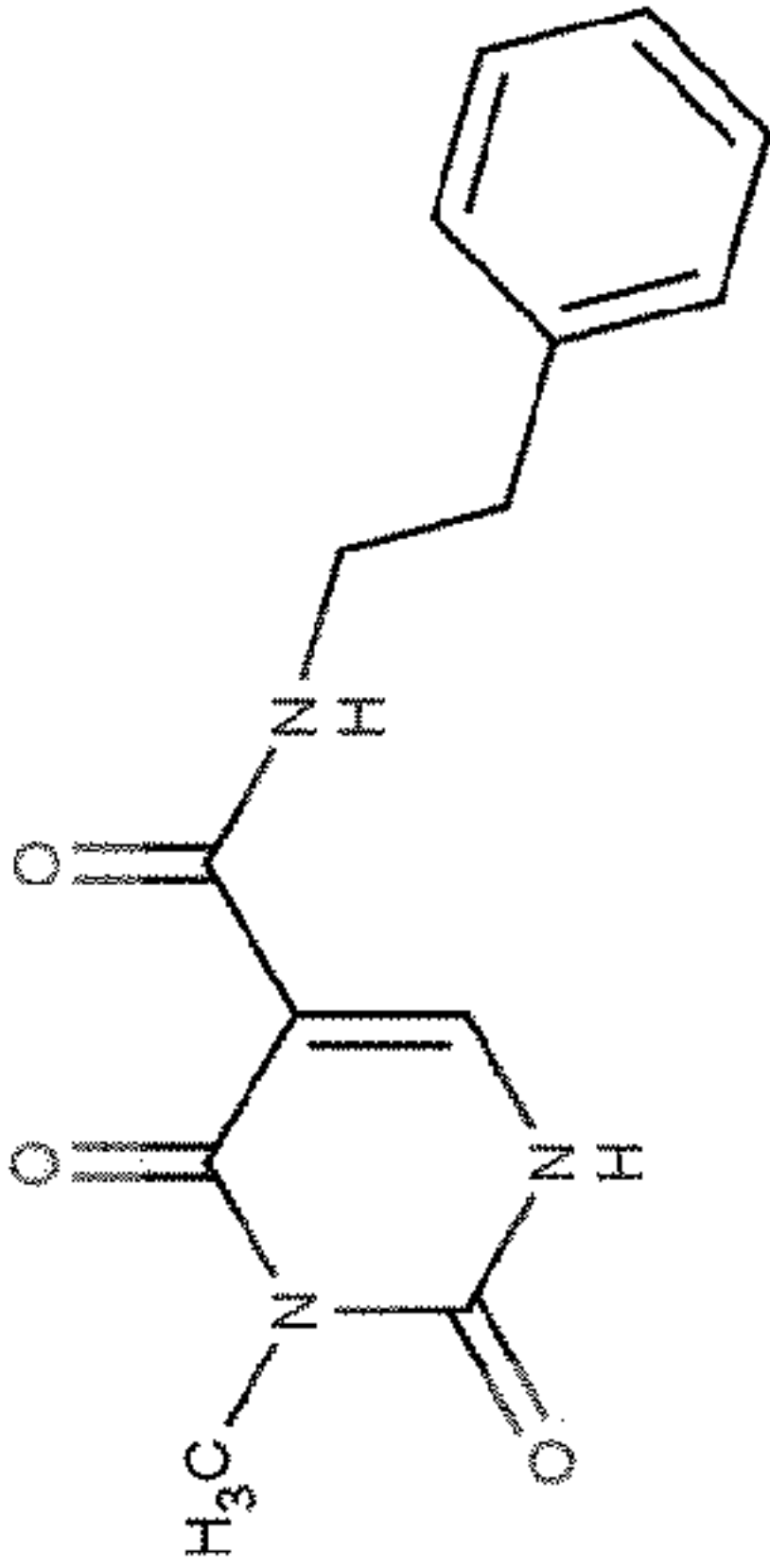
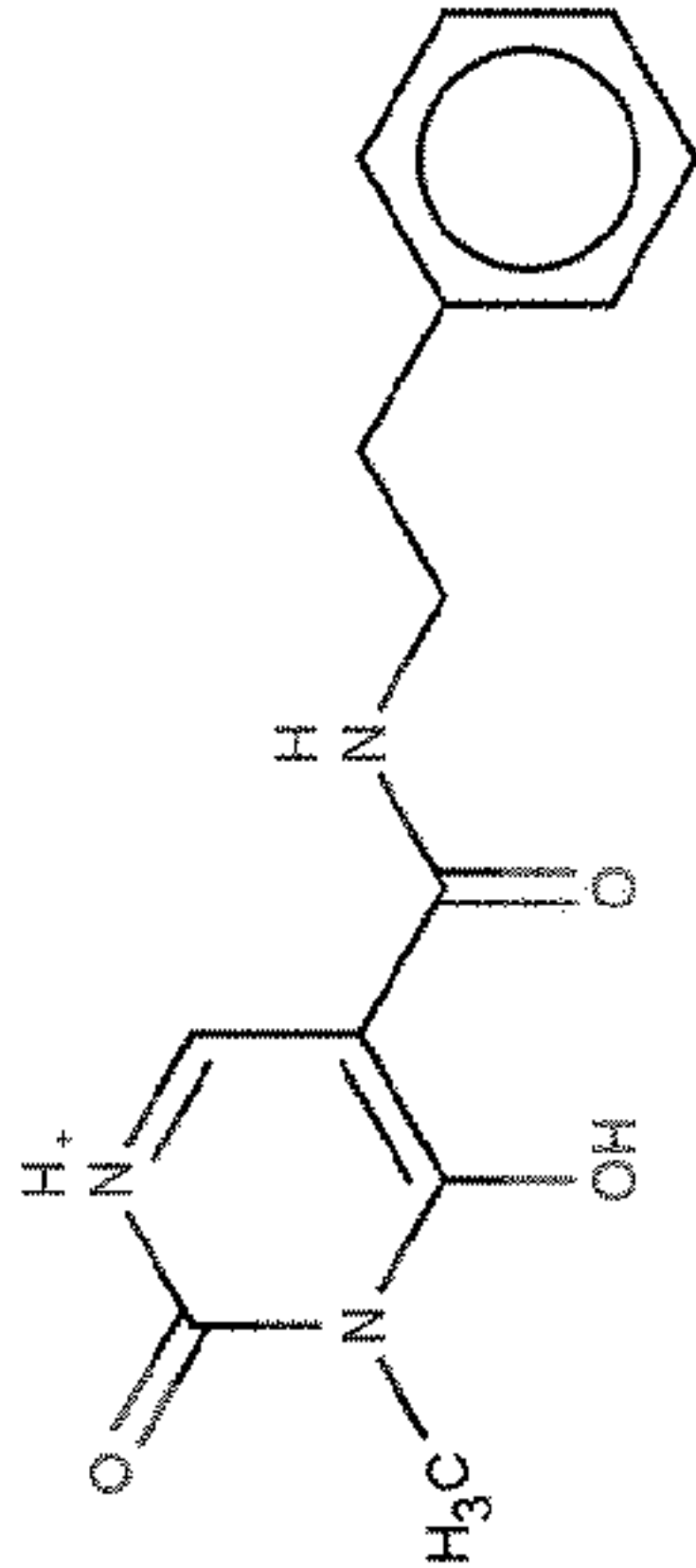
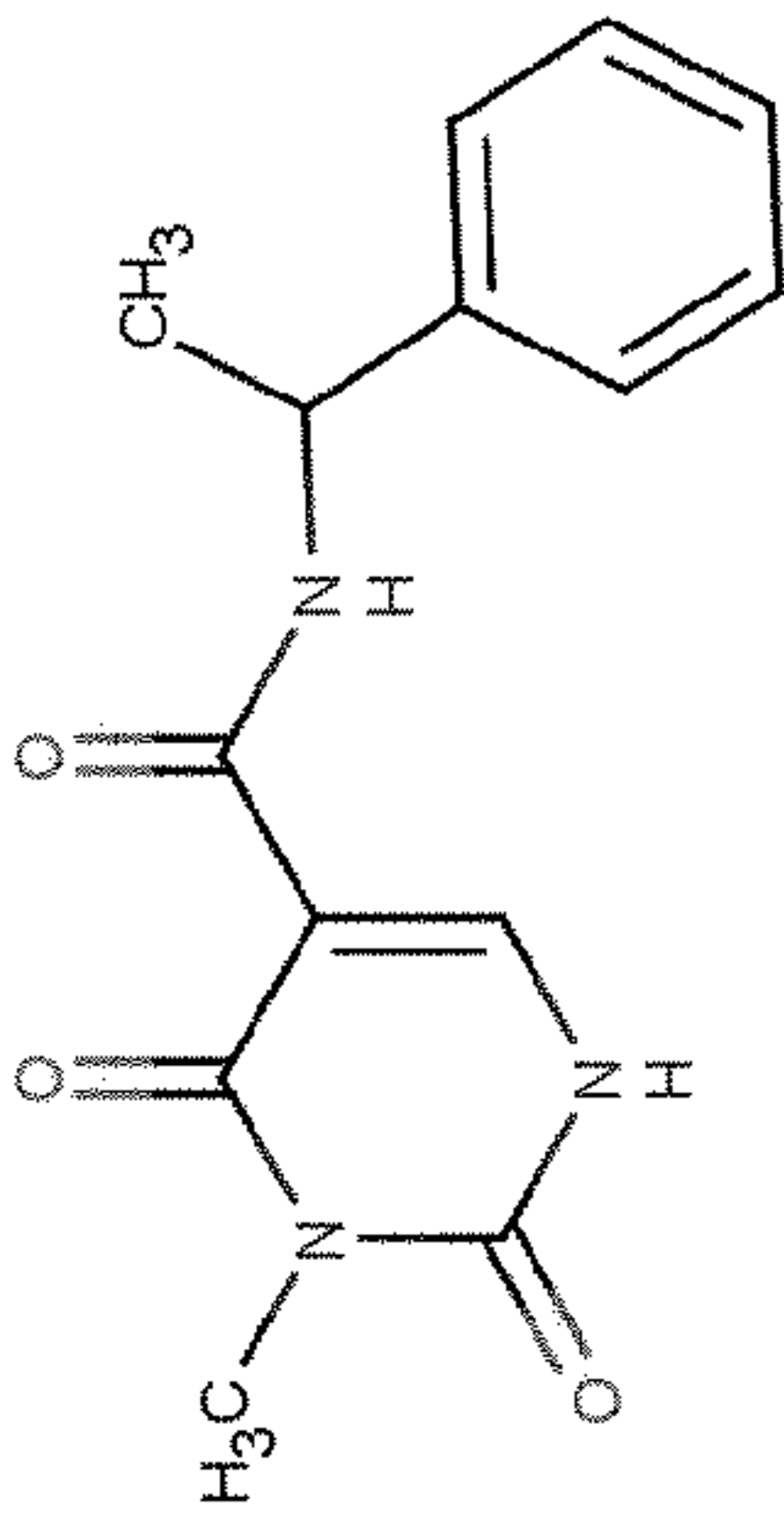
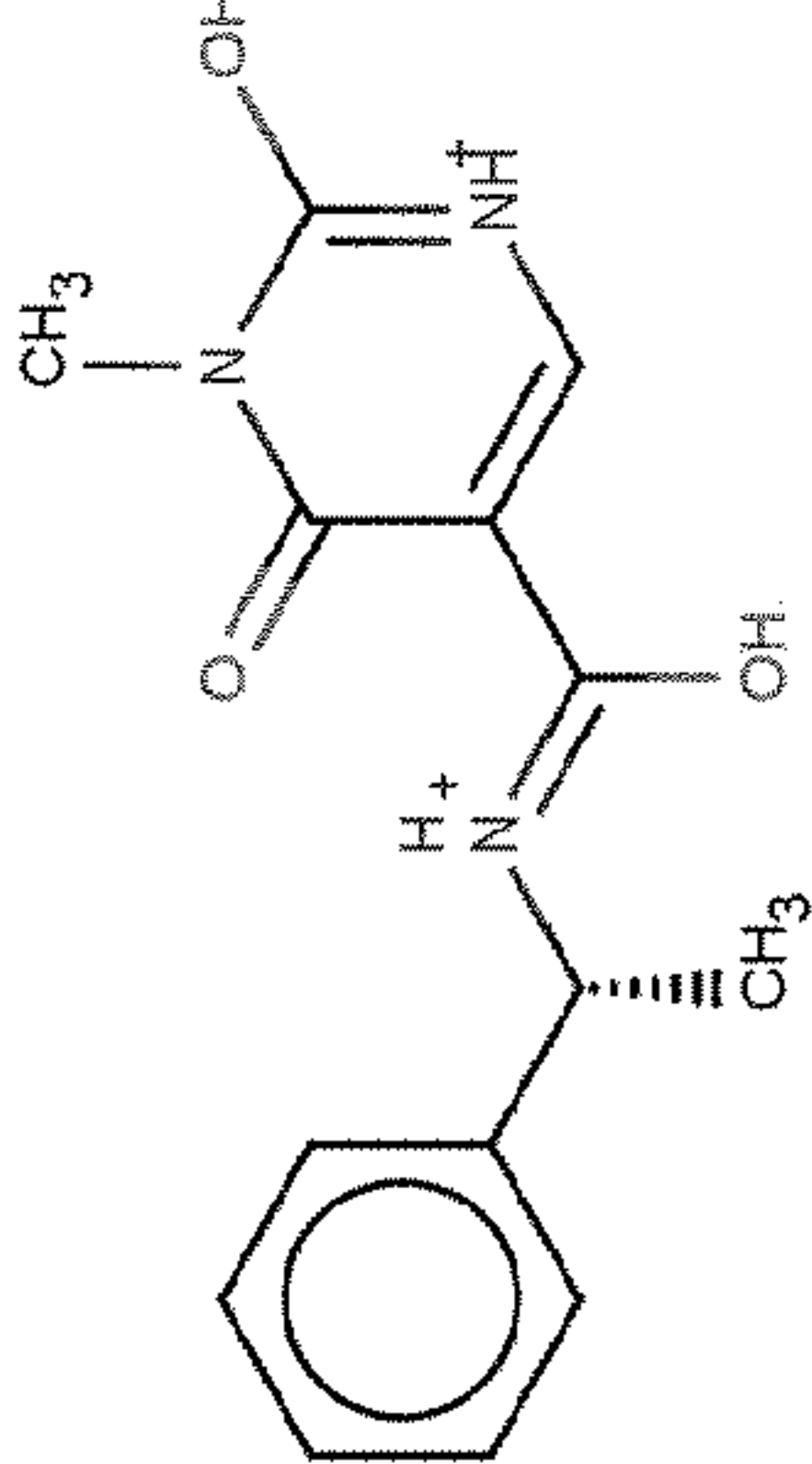
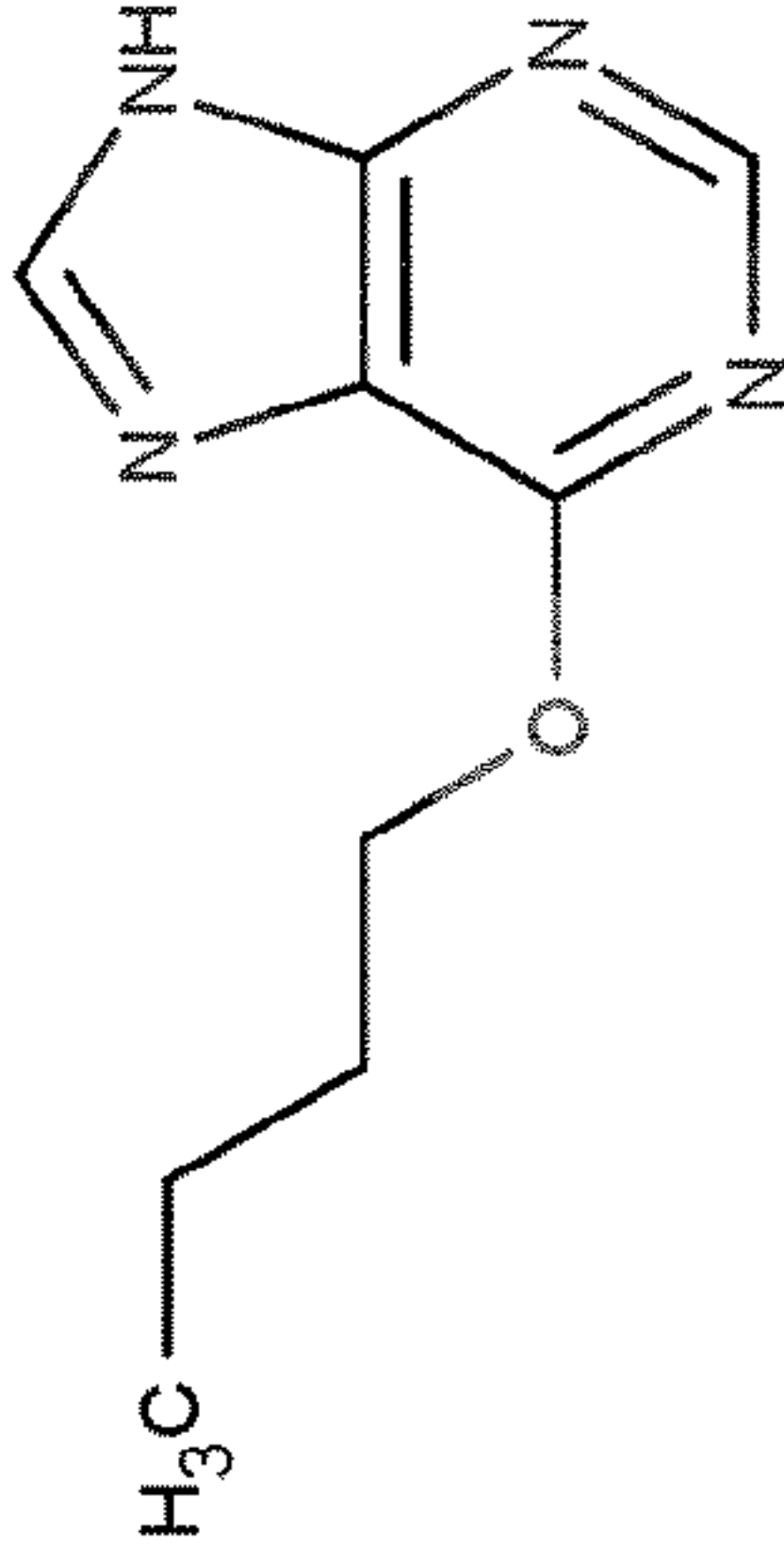
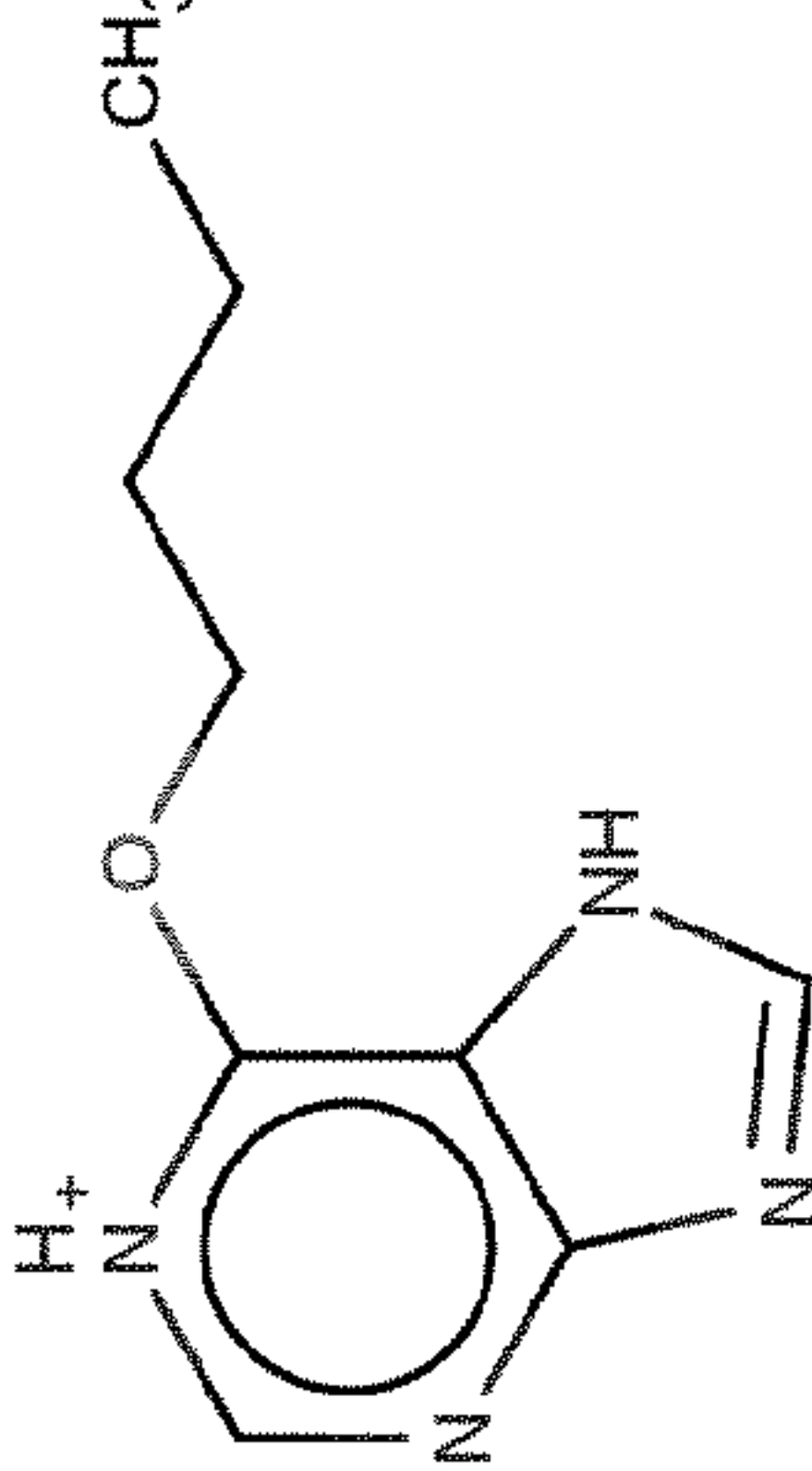
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
	one			
Compound 5	 Preferred IUPAC Name = 5-methoxy-2-(piperazin-1-ylmethyl)-1H-1,3-Benzodiazole	F2145-0409		F2145-0409_5_1 (1 of 14)
Compound 6		F2622-0002		F2622-0002_7_1 (1 of 40)

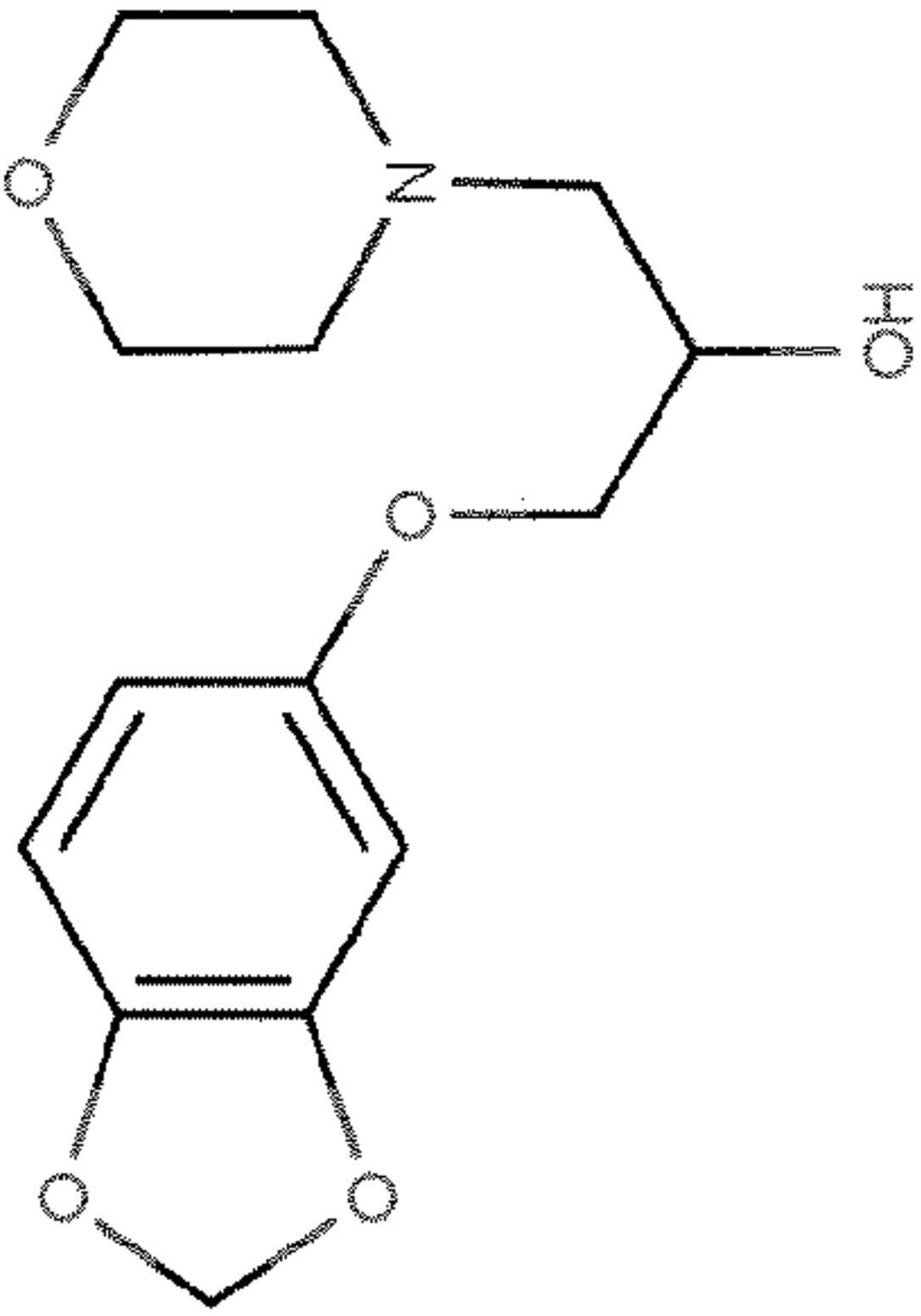
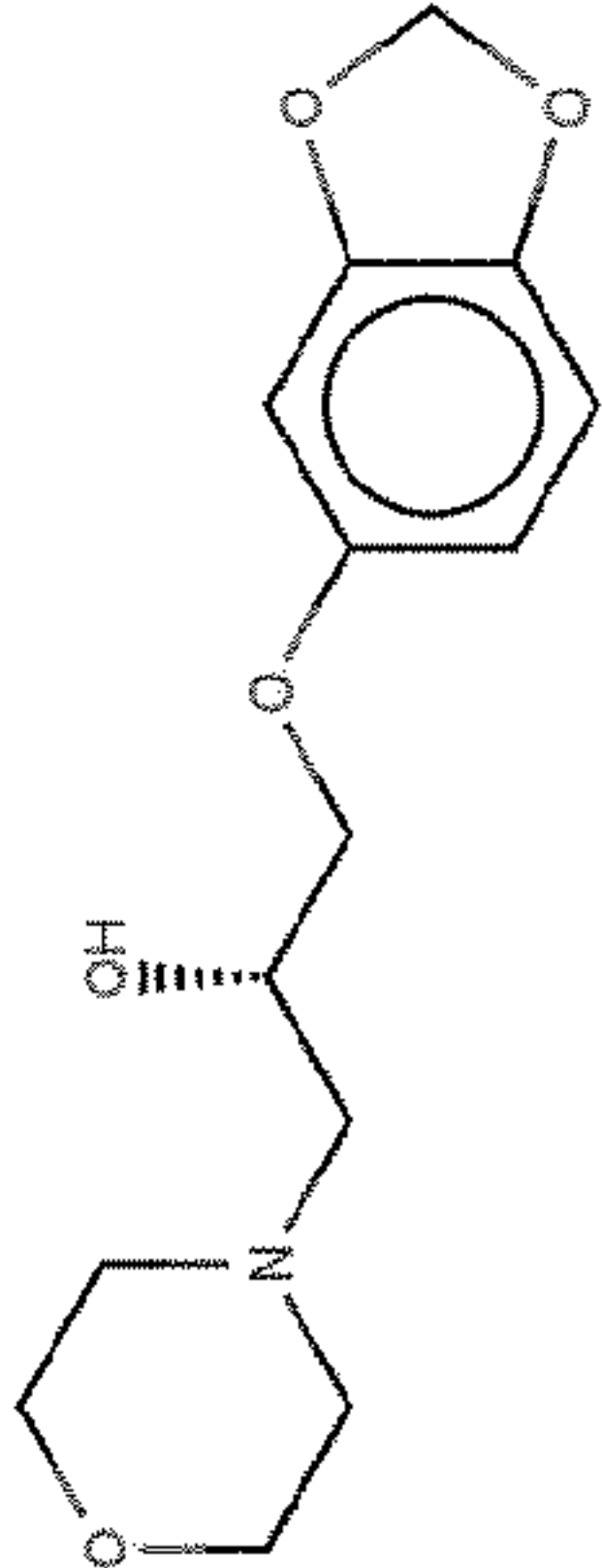
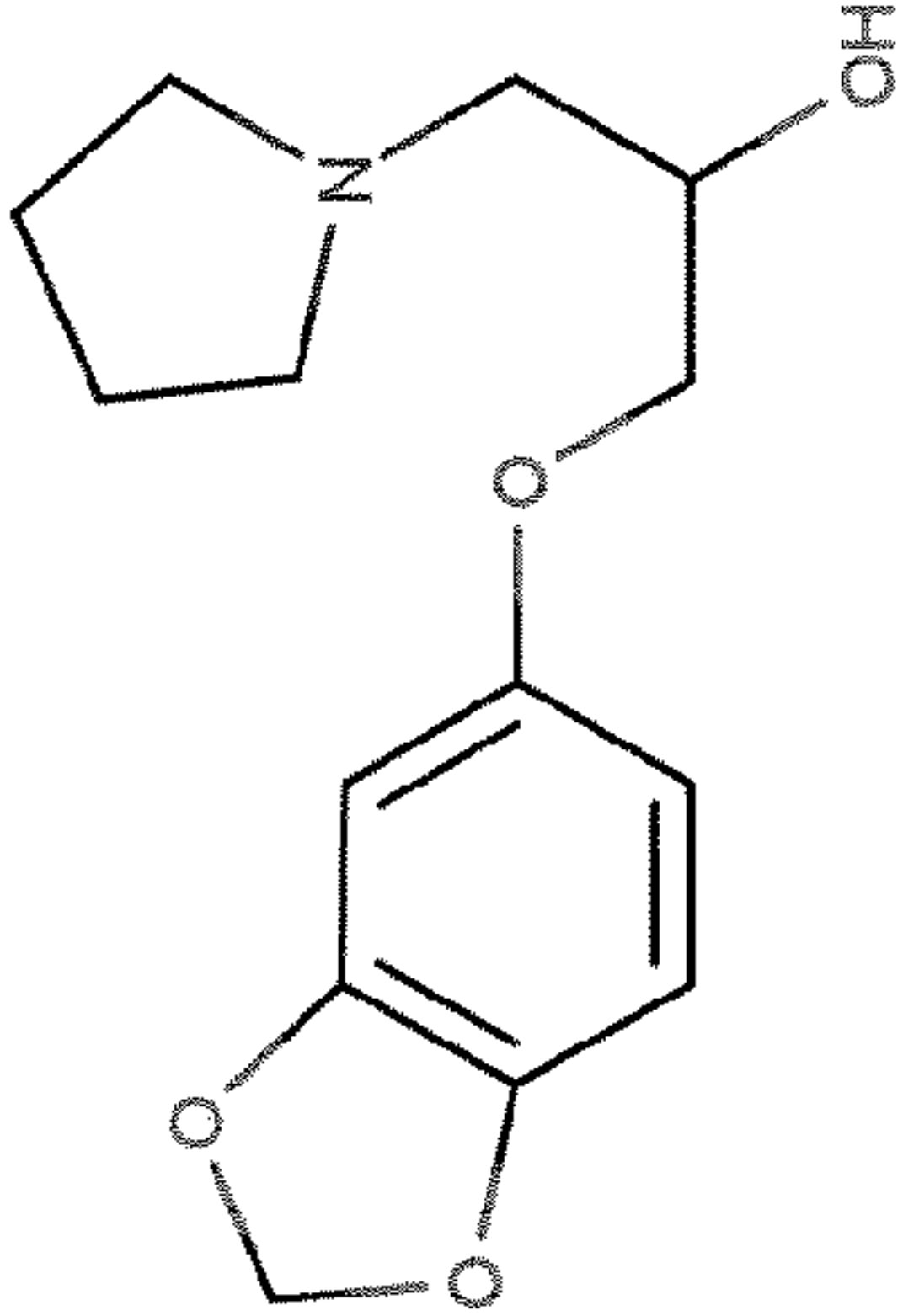
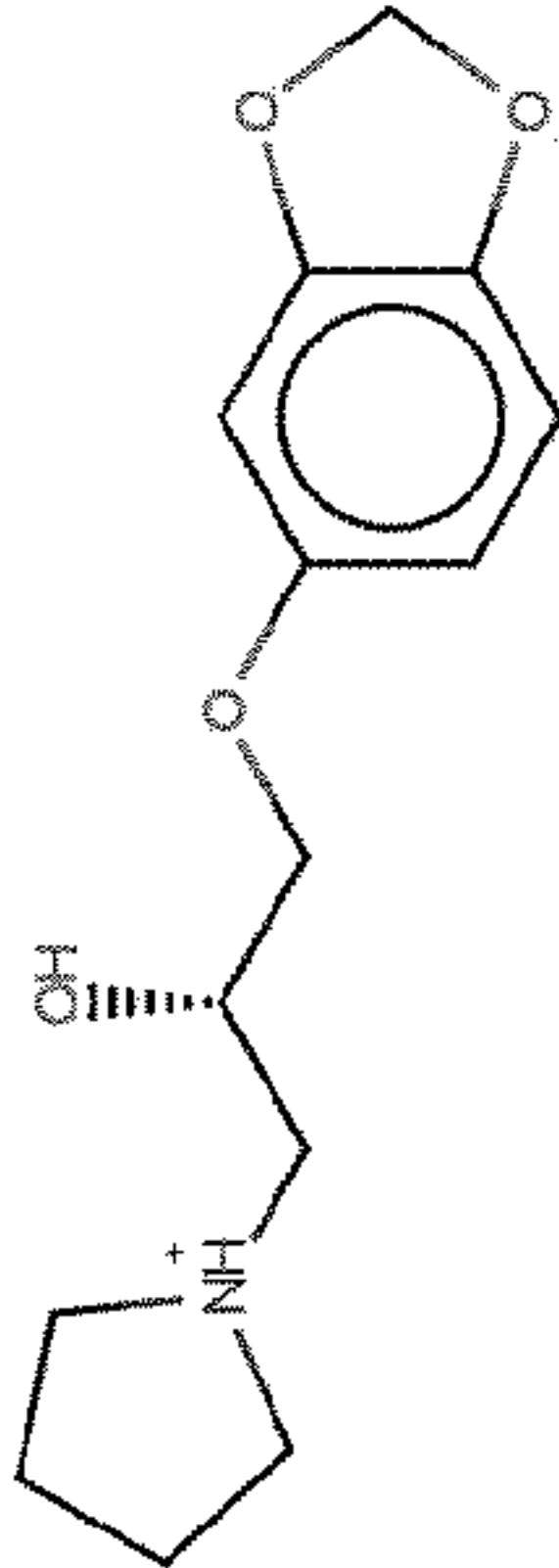
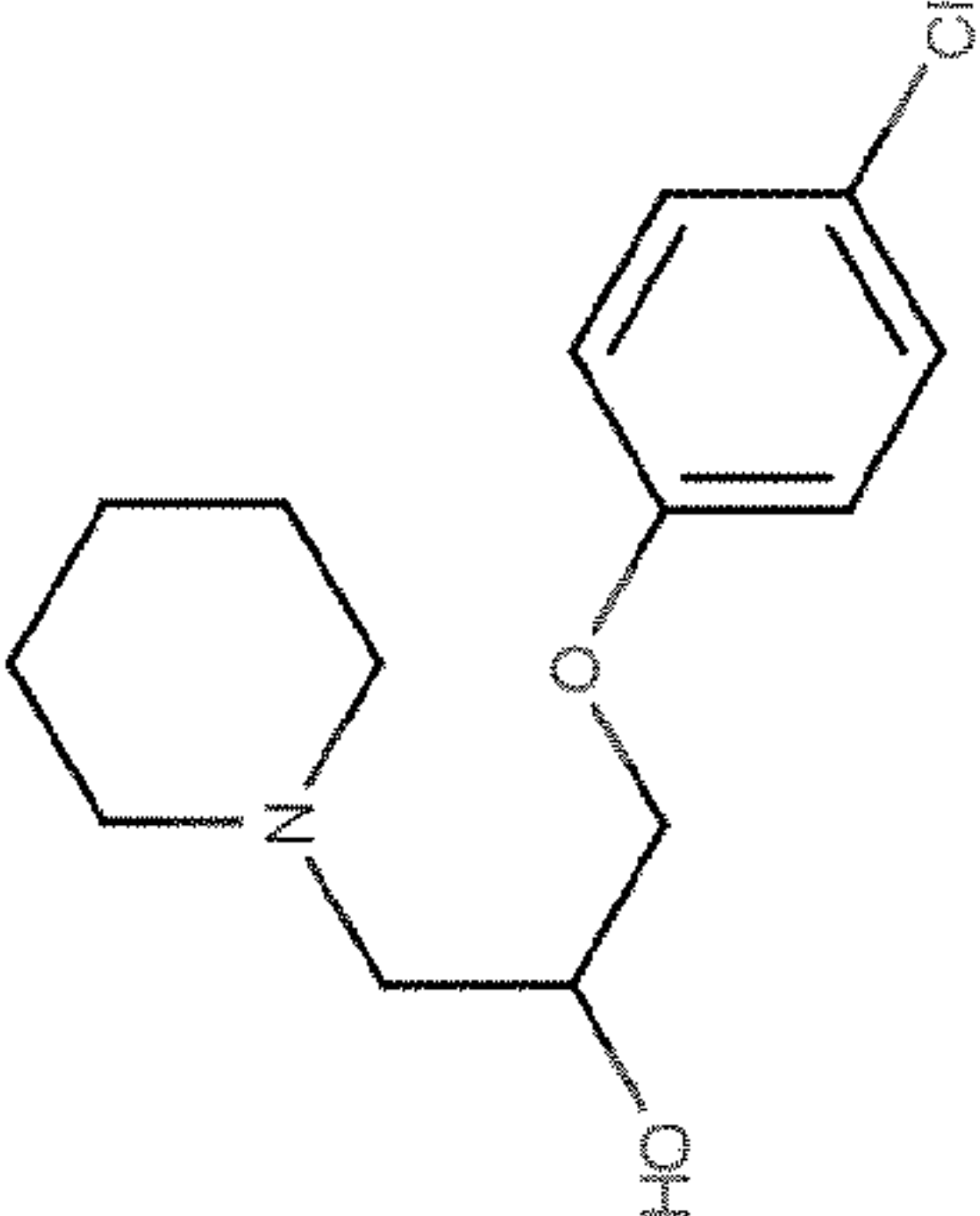
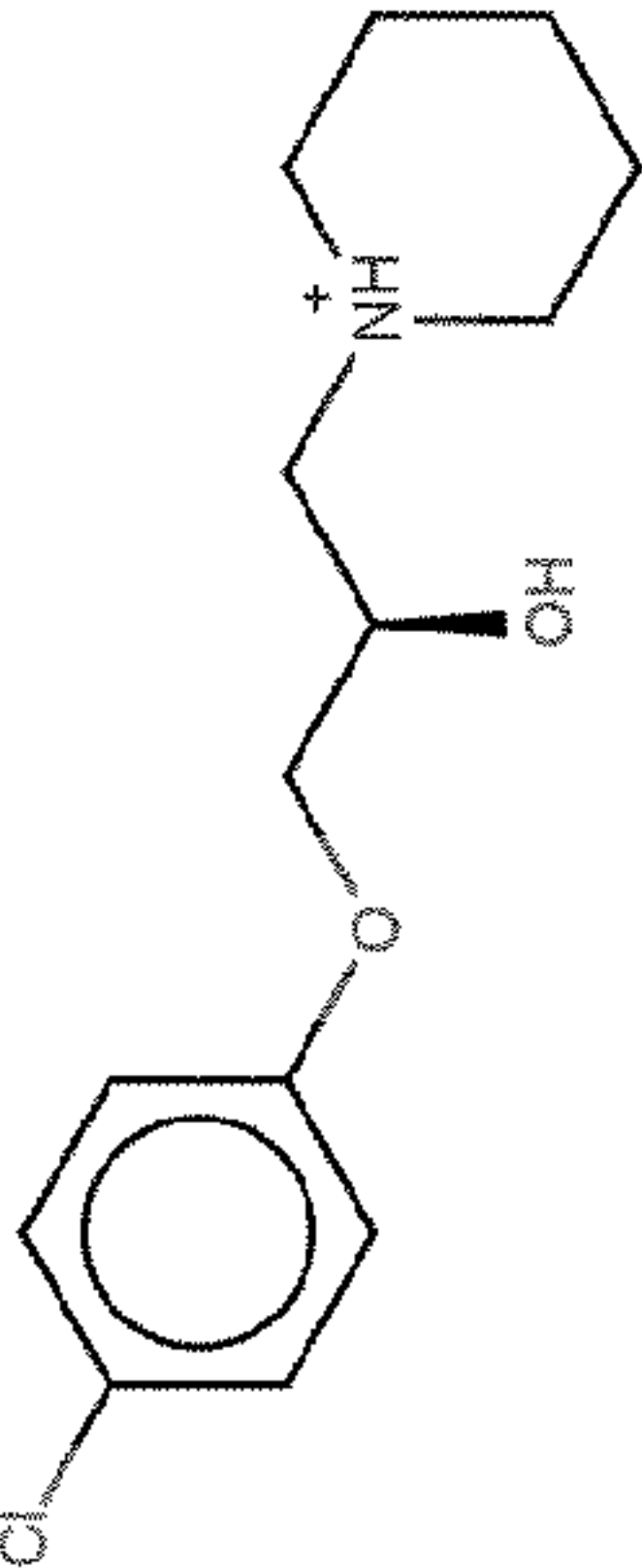
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
	Preferred IUPAC Name = N-butyl-2-hydroxy-9-methyl-4-oxo-4H-pyrido[1,2-a]pyrimidine-3-carboxamide			
Compound 7		F0050-0006		F0050-0006_2 (1 of 4)
Compound 8		F0095-1317		F0095-1317_13_5 (1 of 116)

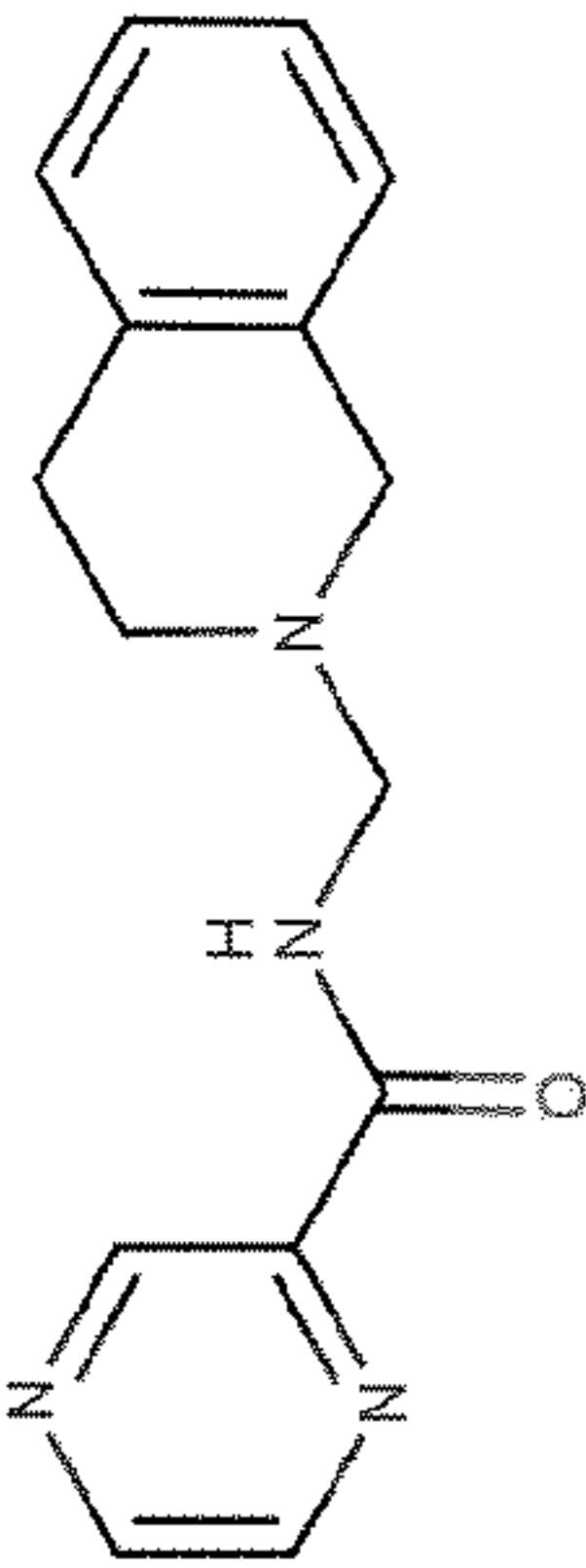
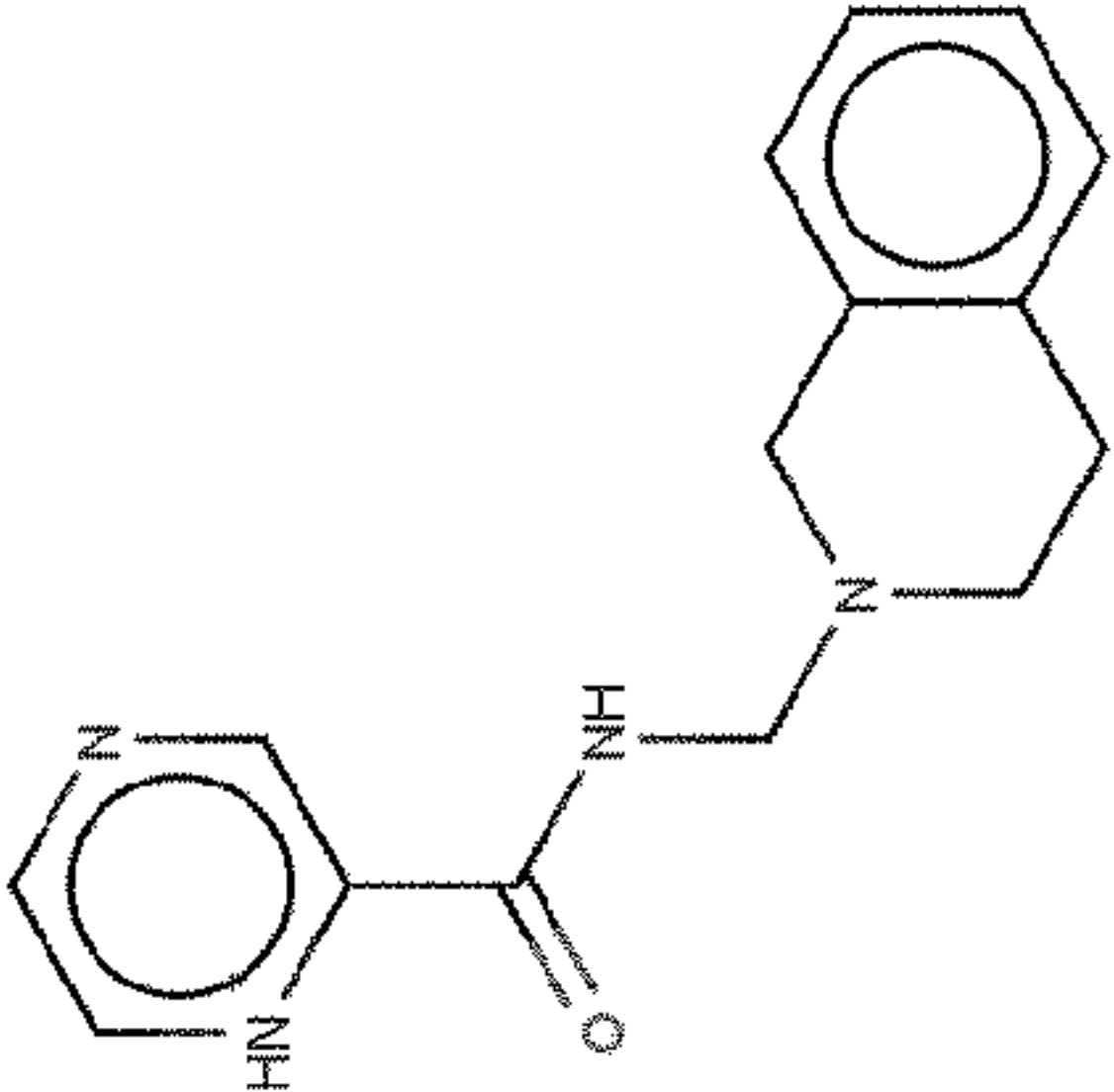
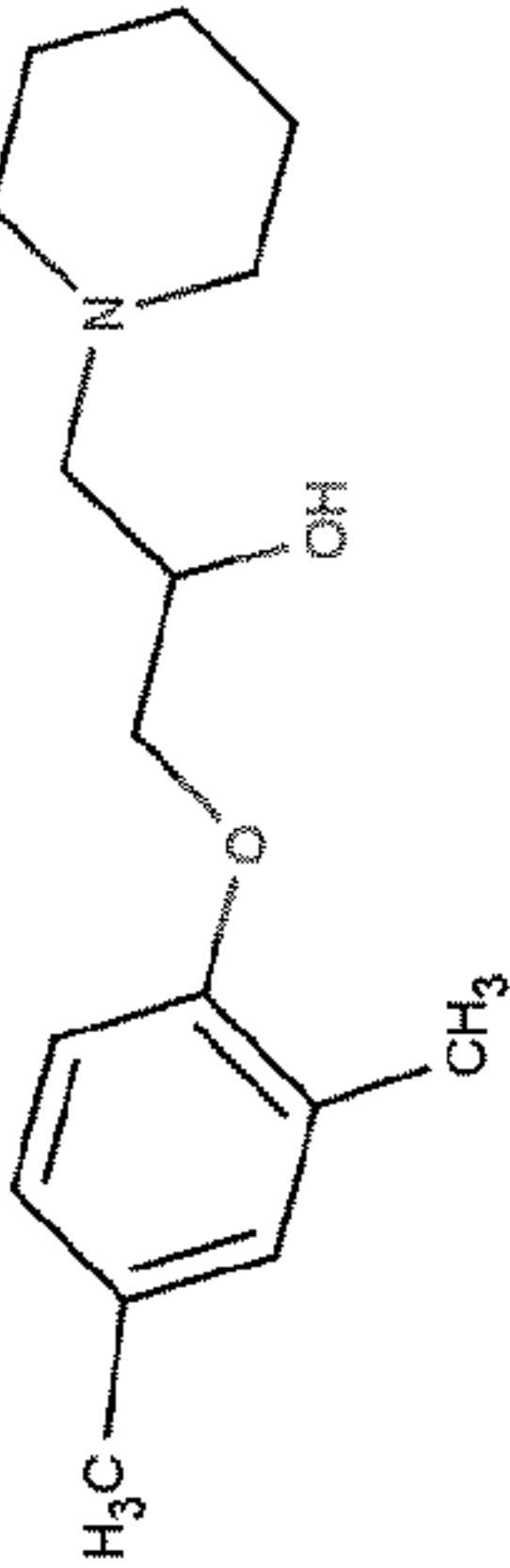
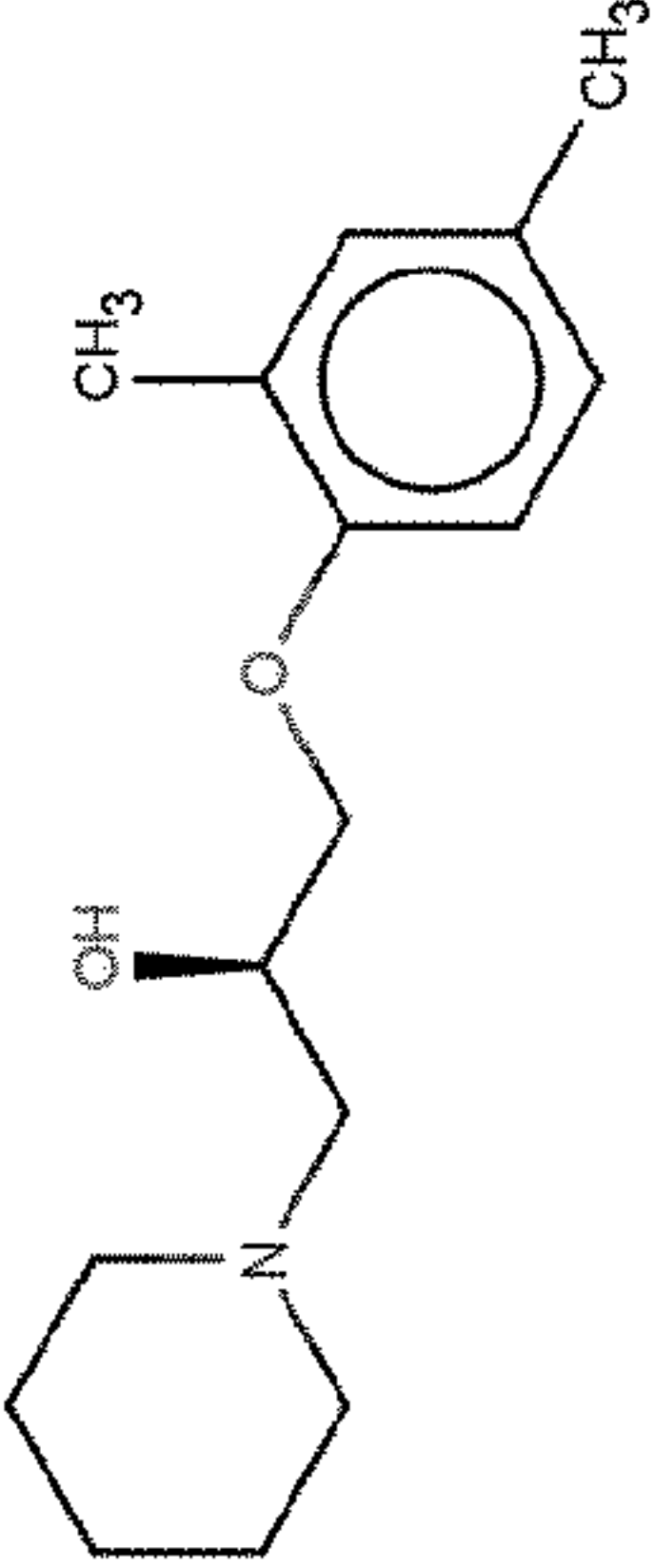
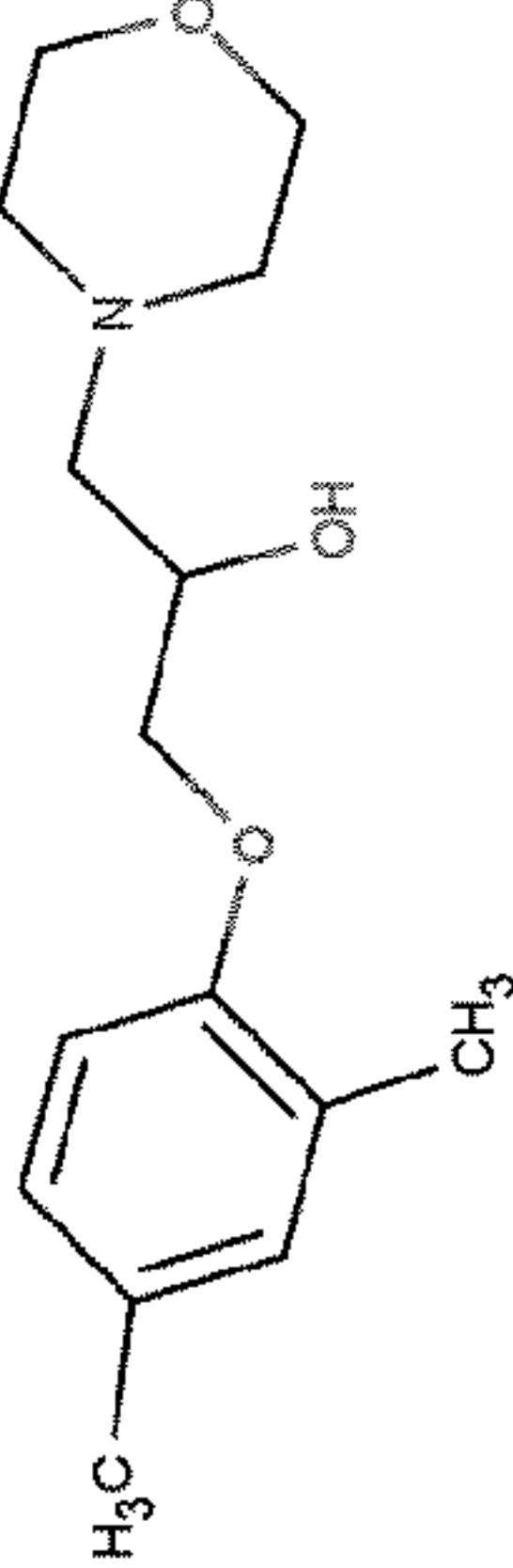
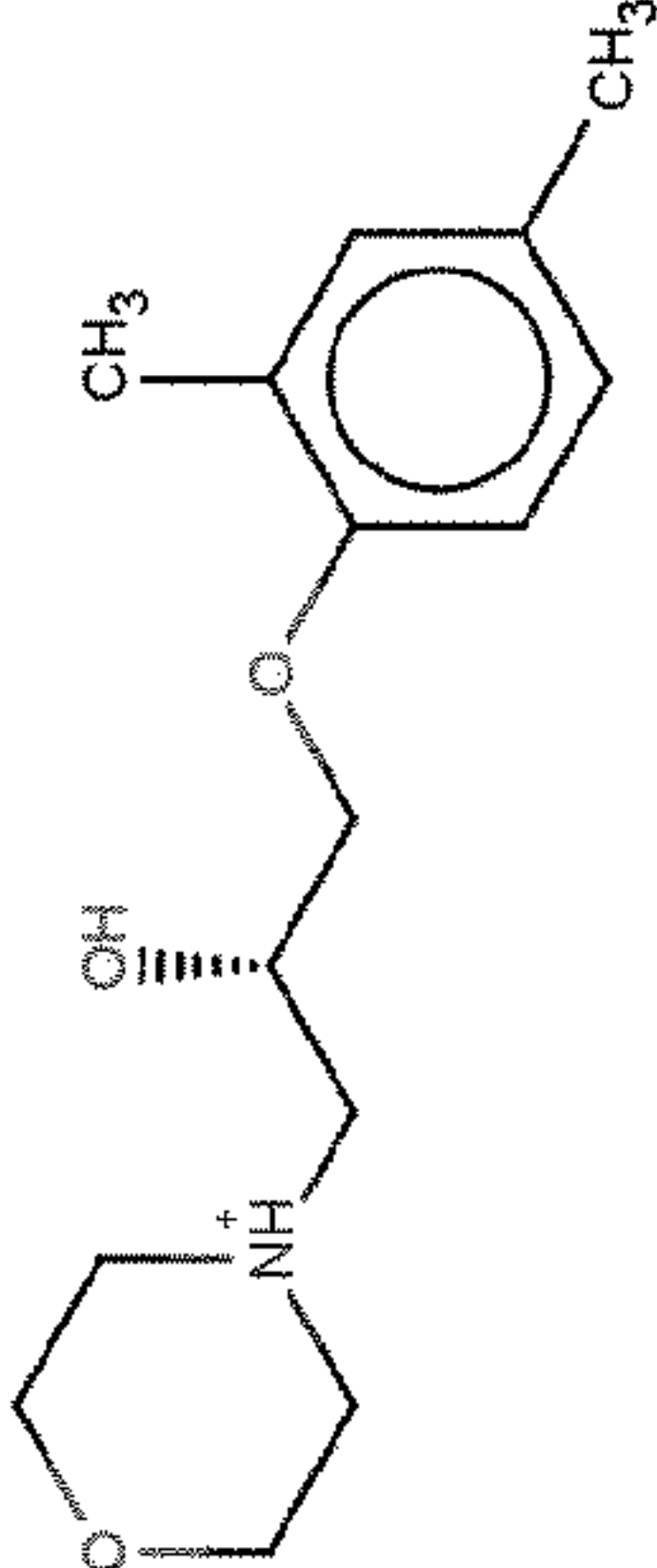
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 9		F0126-0203		F0126-0203_16_4 (1 of 46)
Compound 10		F0176-0083		F0176-0083_3_1 (1 of 10)
Compound 11		F0196-0408		F0196-0408_1_1 (1 of 2)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 12		F0285-0322		F0285-0322_2 (1 of 2)
Compound 13		F0347-0341		F0347-0341_21_1 (1 of 14)
Compound 14		F0347-0841		F0347-0841_18_1 (1 of 12)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 15		F0578-0190		F0578-0190_7_1 (1 of 22)
Compound 16		F0611-0055		F0611-0055_1 (1 of 4)
Compound 17		F0693-0722		F0693-0722_14_1 (1 of 22)

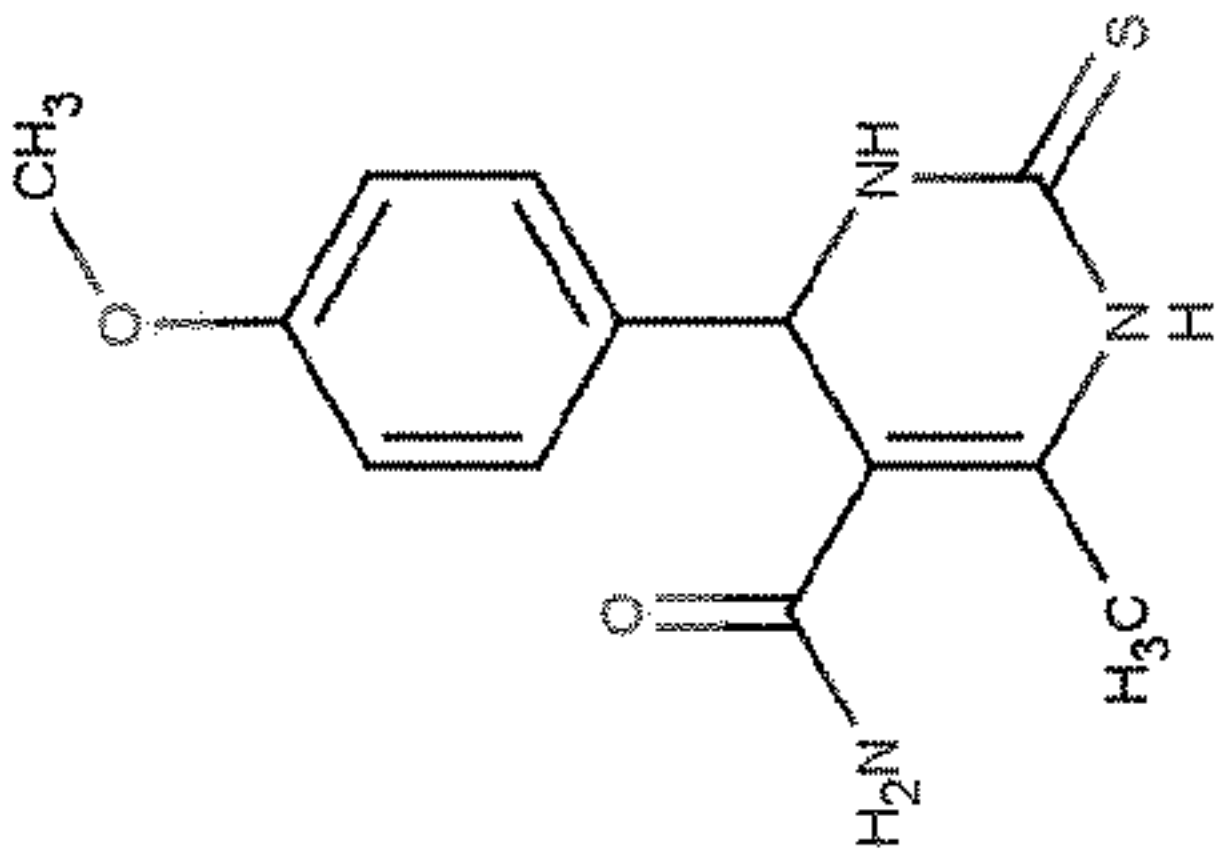
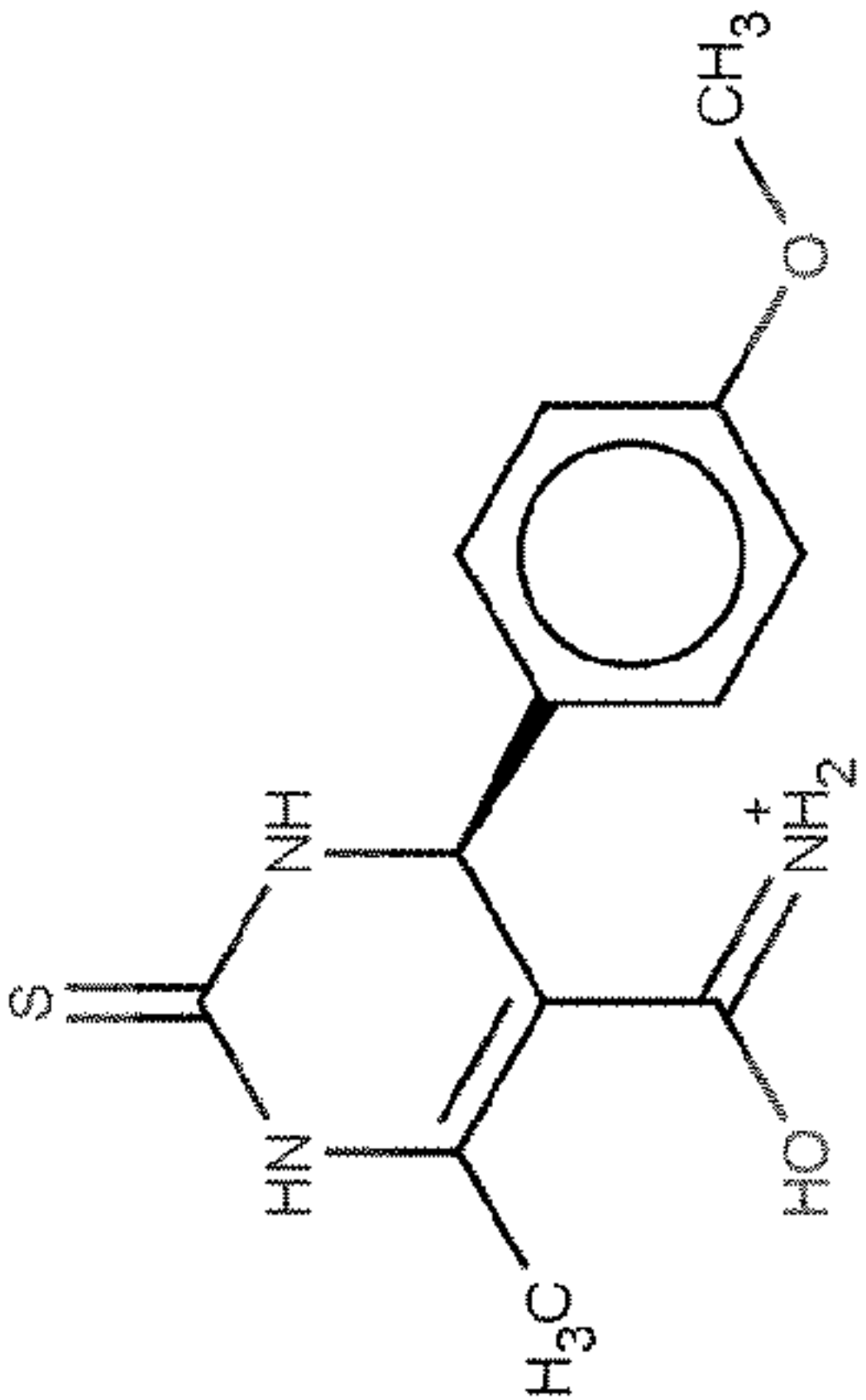
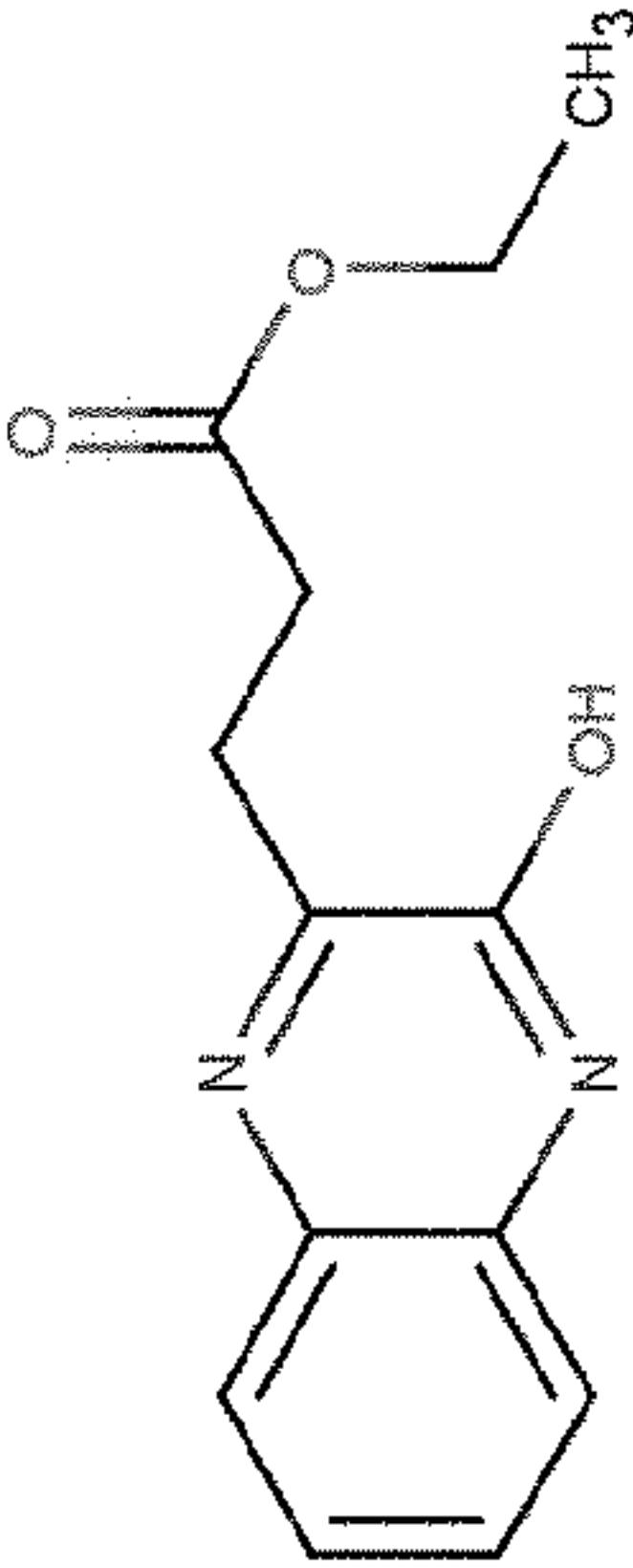
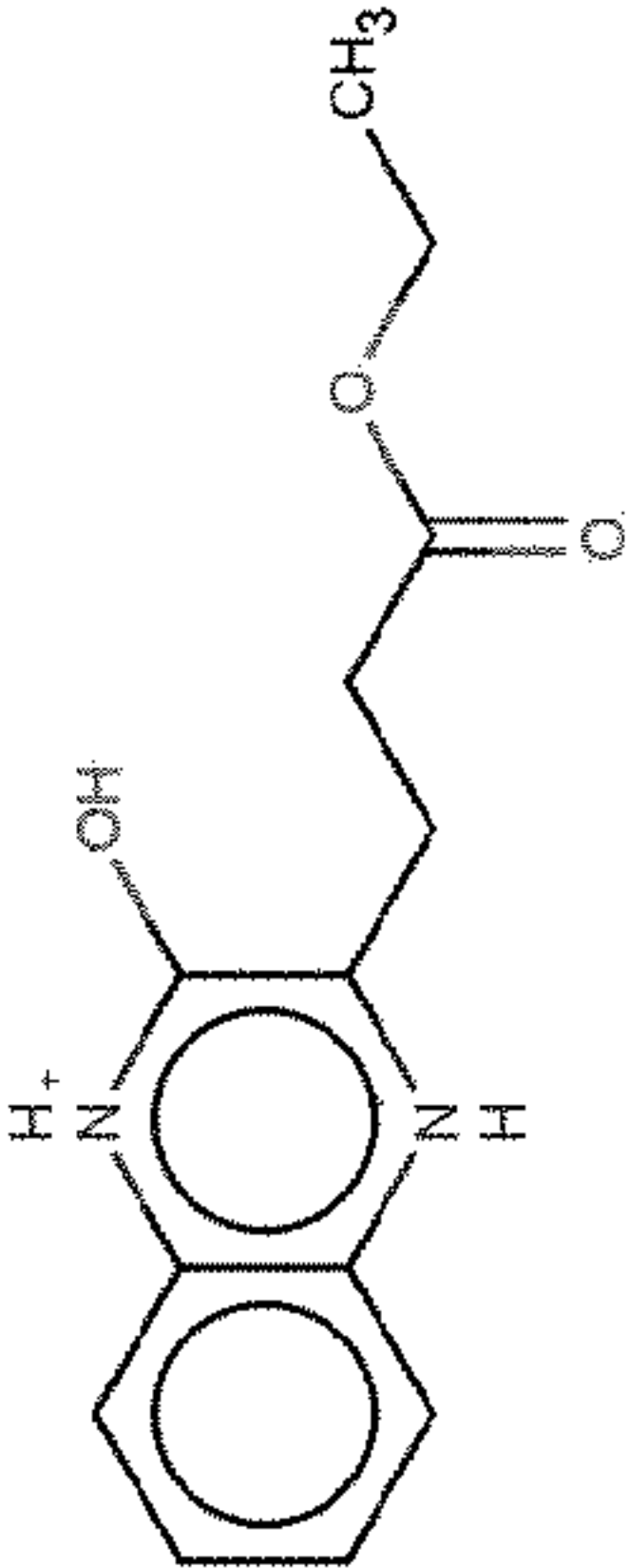
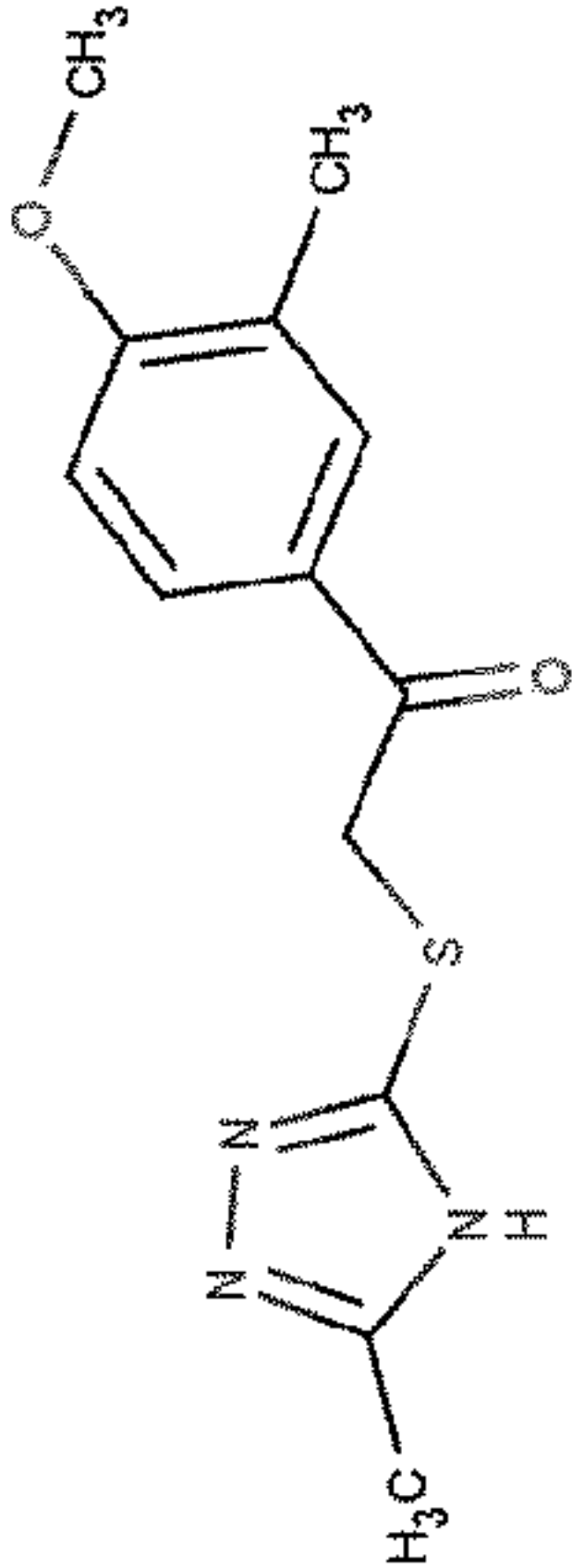
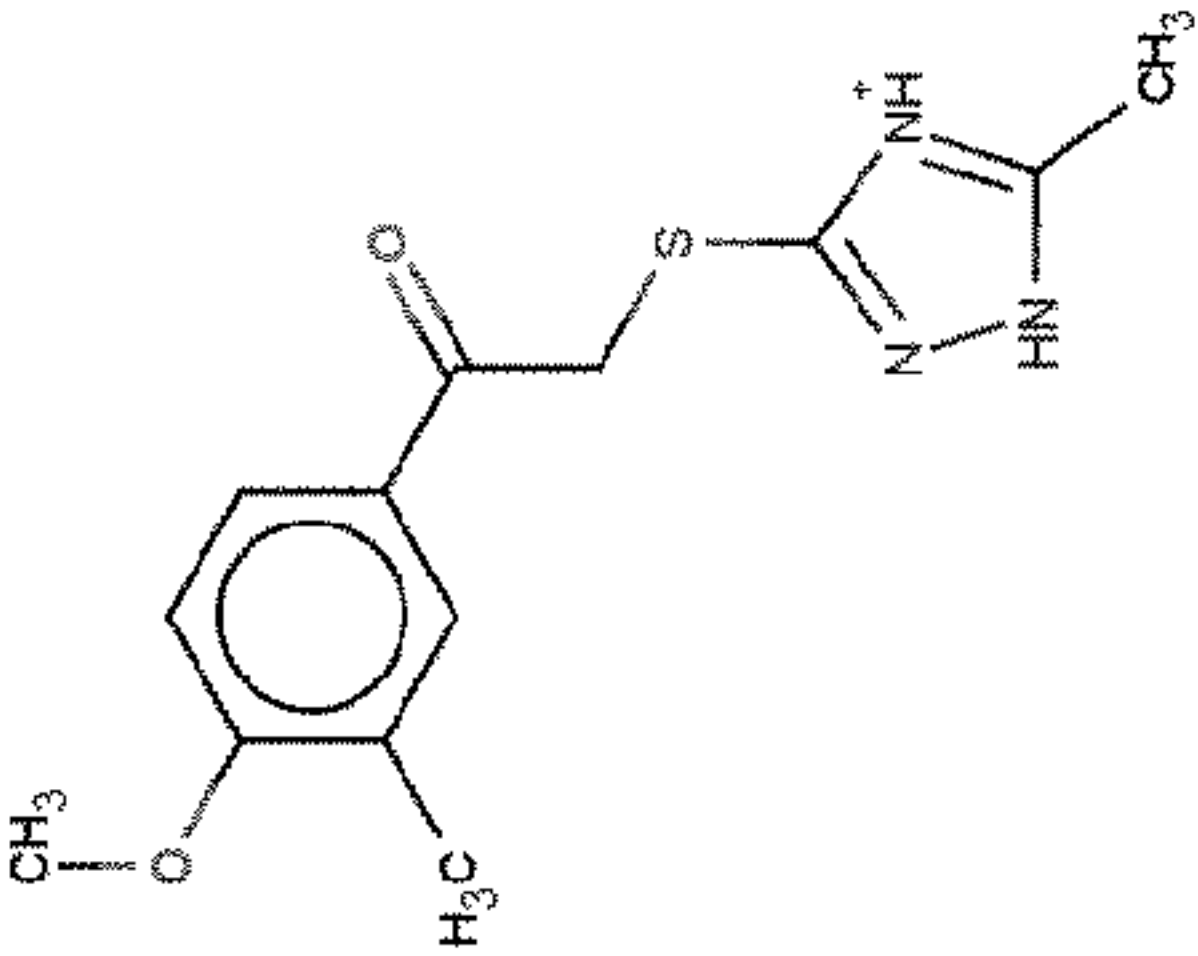
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 18		F0696-0806		F0696-0806_6_1 (1 of 44)
Compound 19		F0696-0849		F0696-0849_24_1 (1 of 88)
Compound 20		F0704-0019		F0704-0019_7_1 (1 of 11)

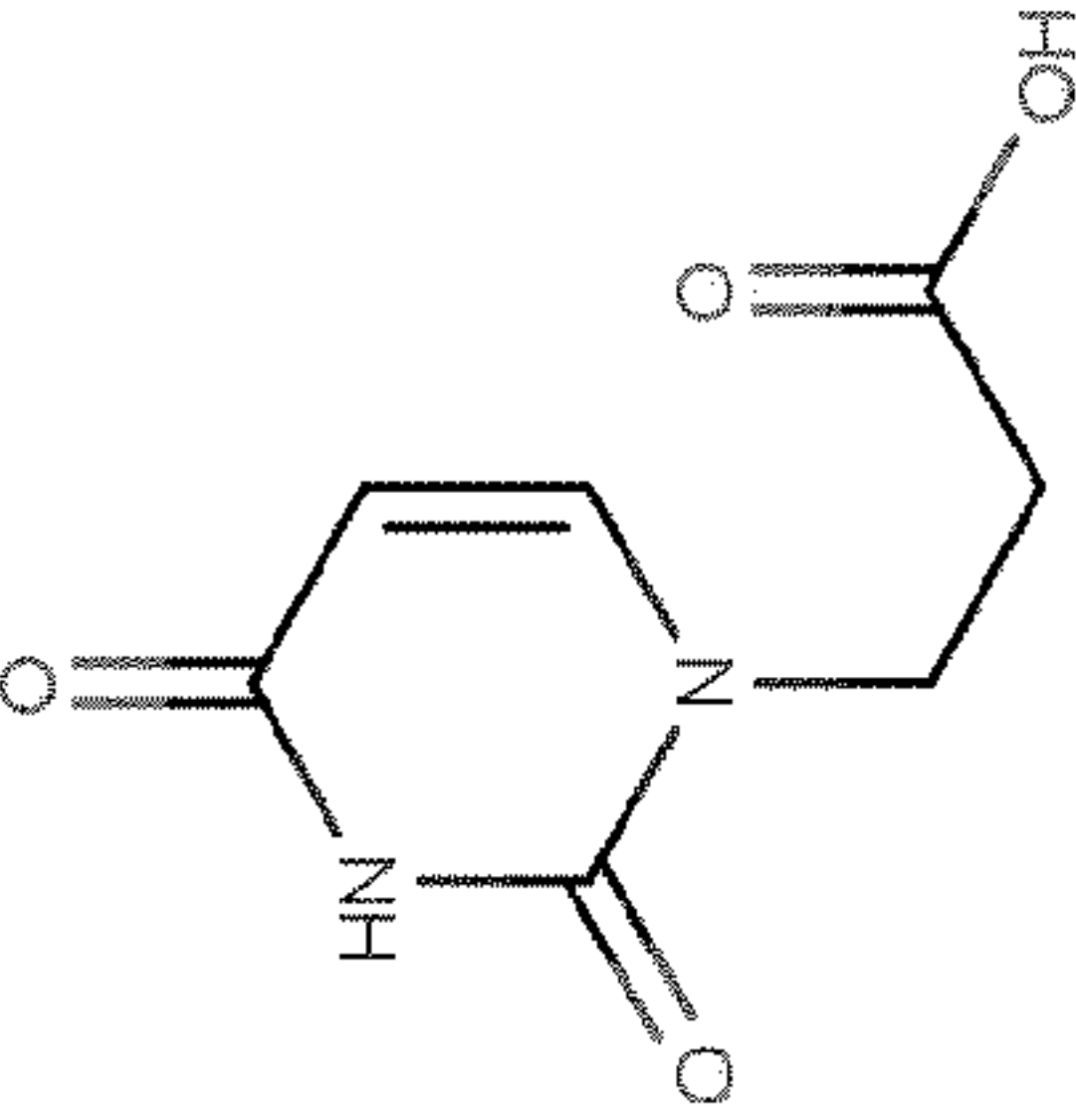
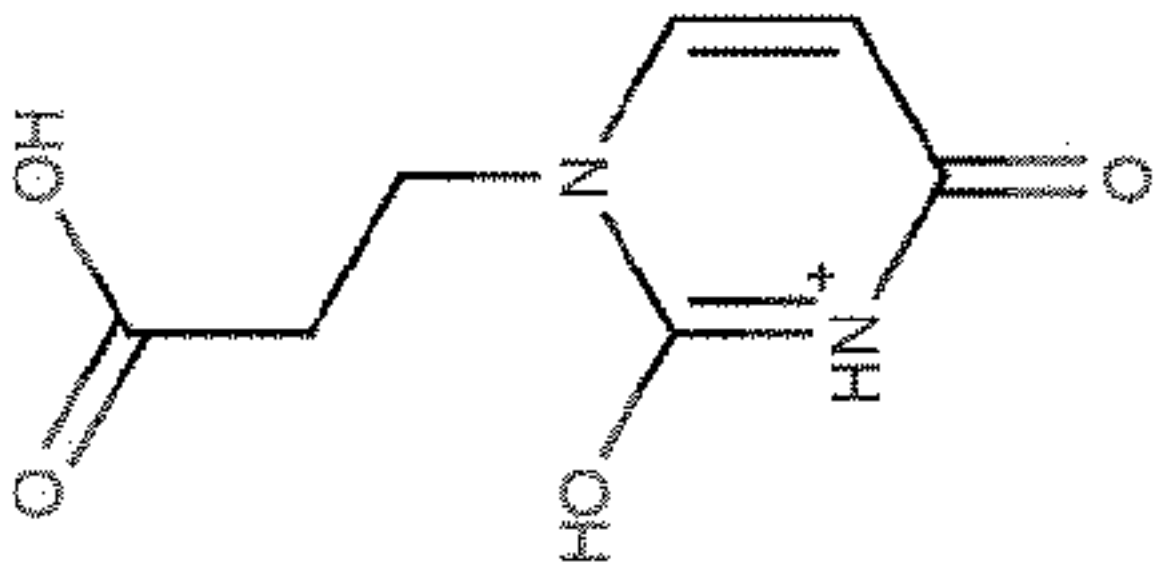
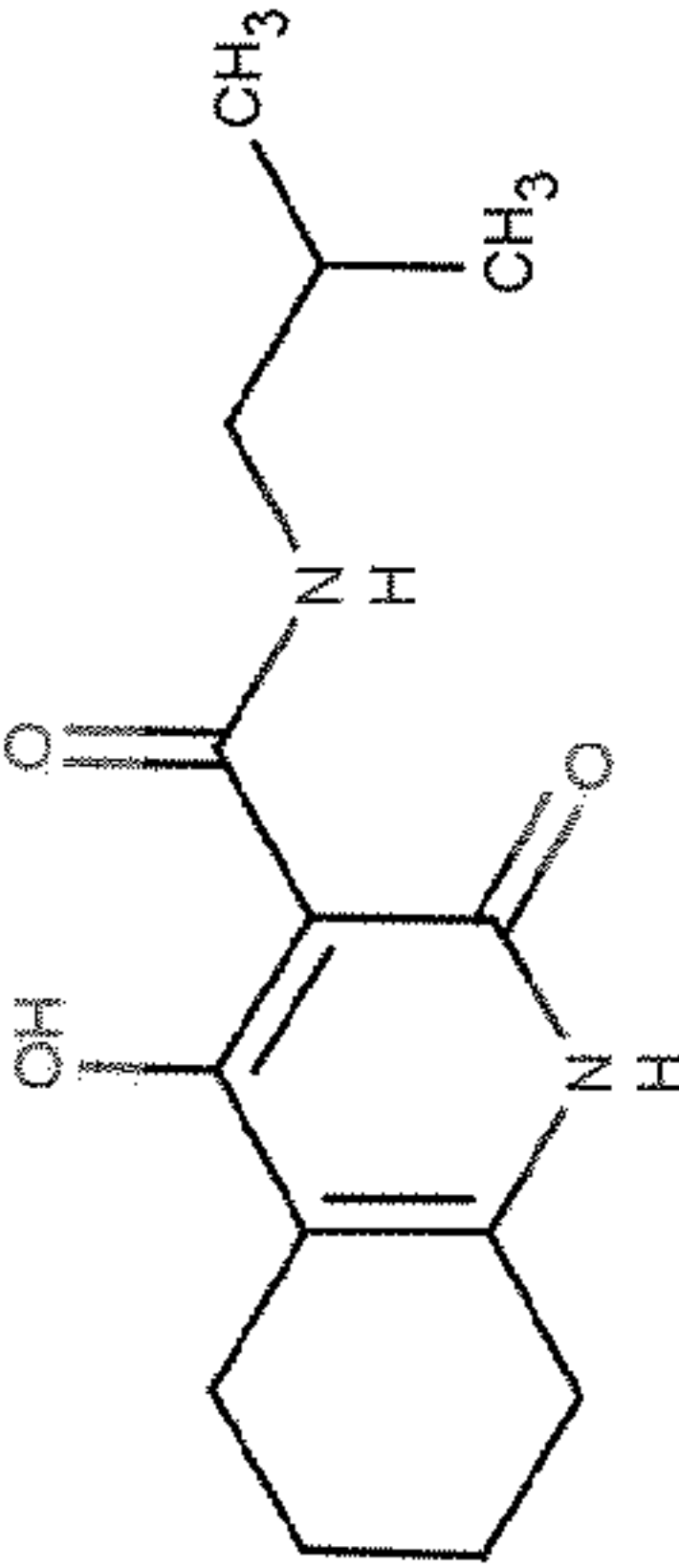
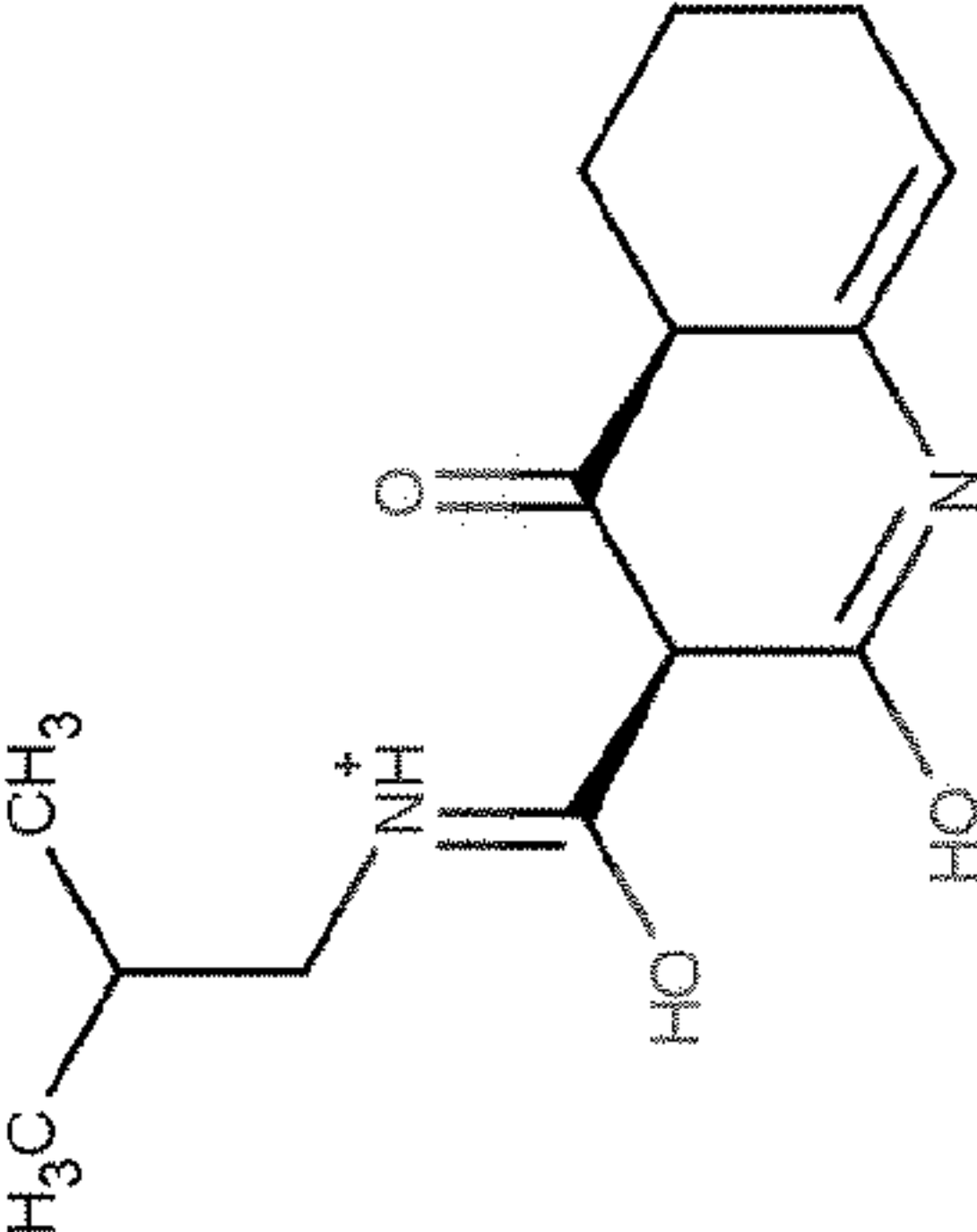
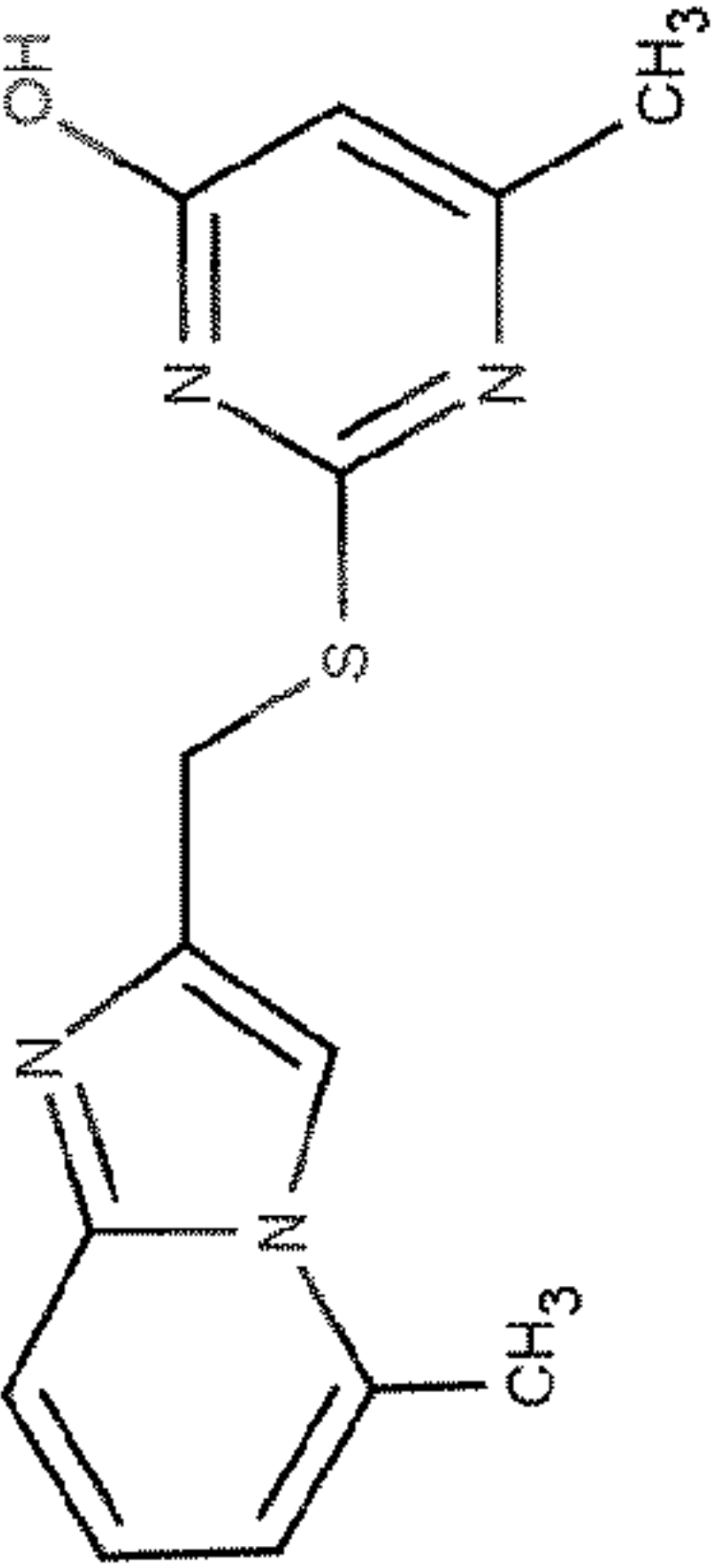
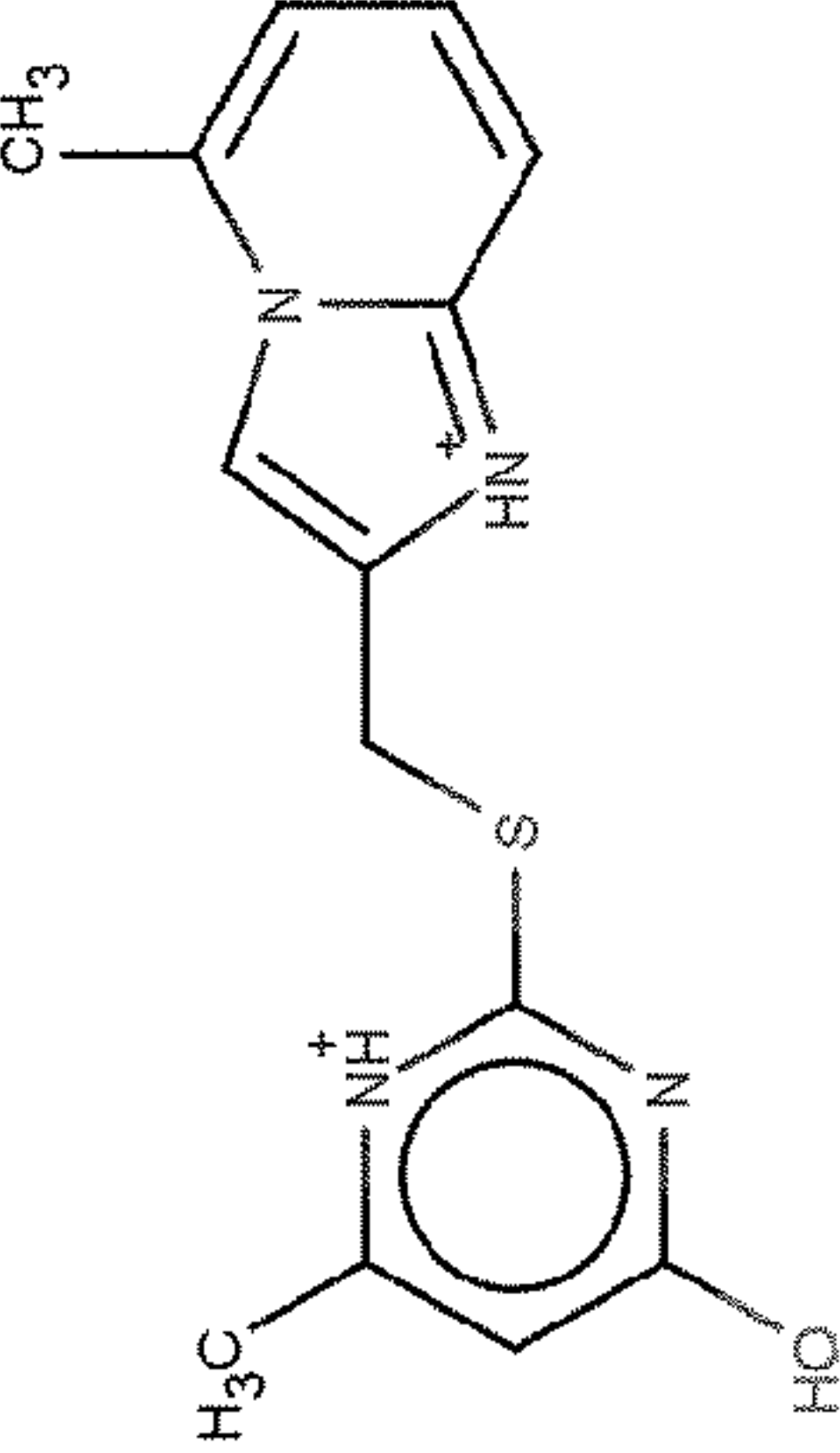
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 21		F0719-00666		F0719-00666_1 (1 of 4)
Compound 22		F0719-0070		F0719-0070_1_1 (1 of 4)
Compound 23		F0719-0145		F0719-0145_1_1 (1 of 4)

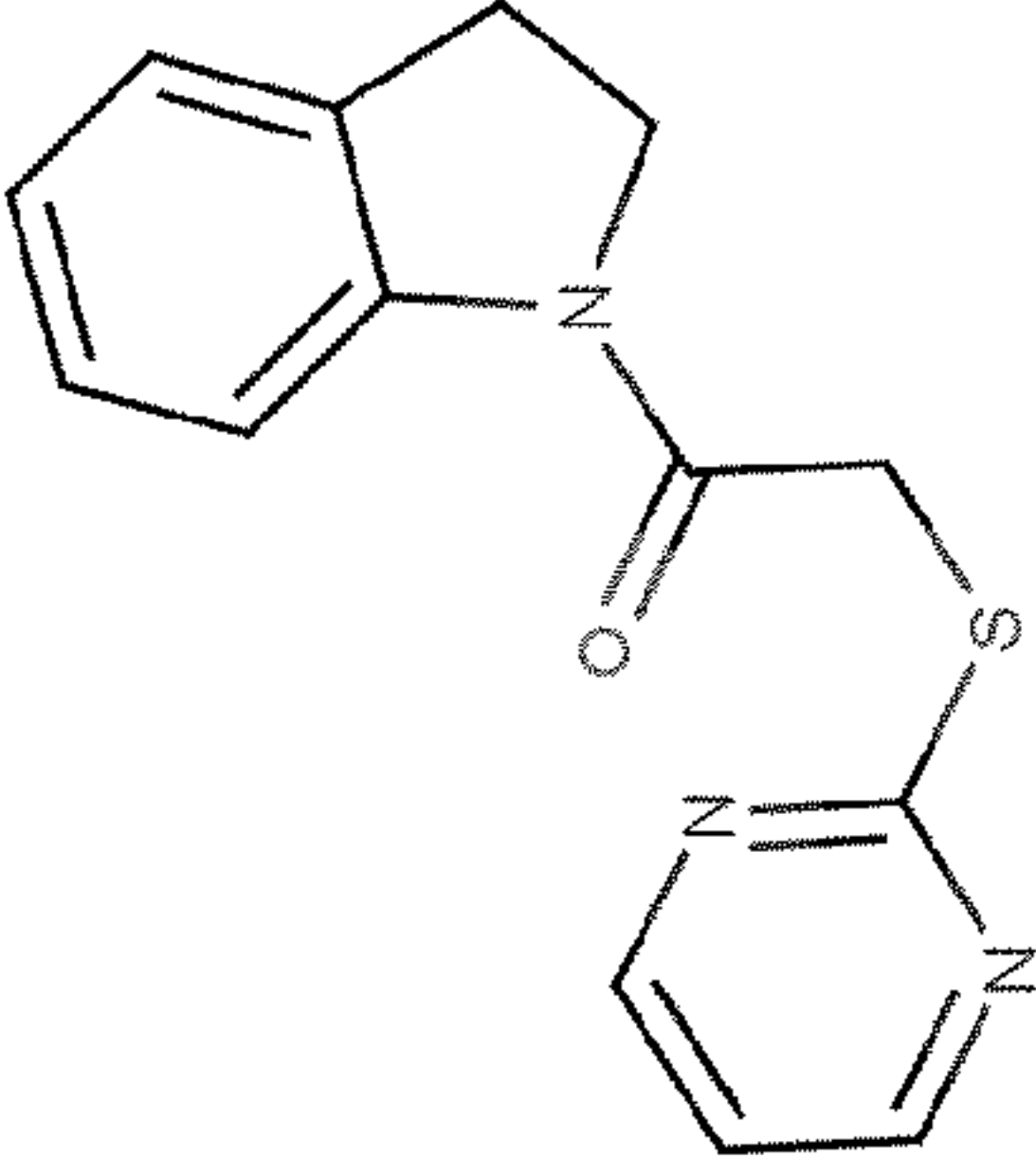
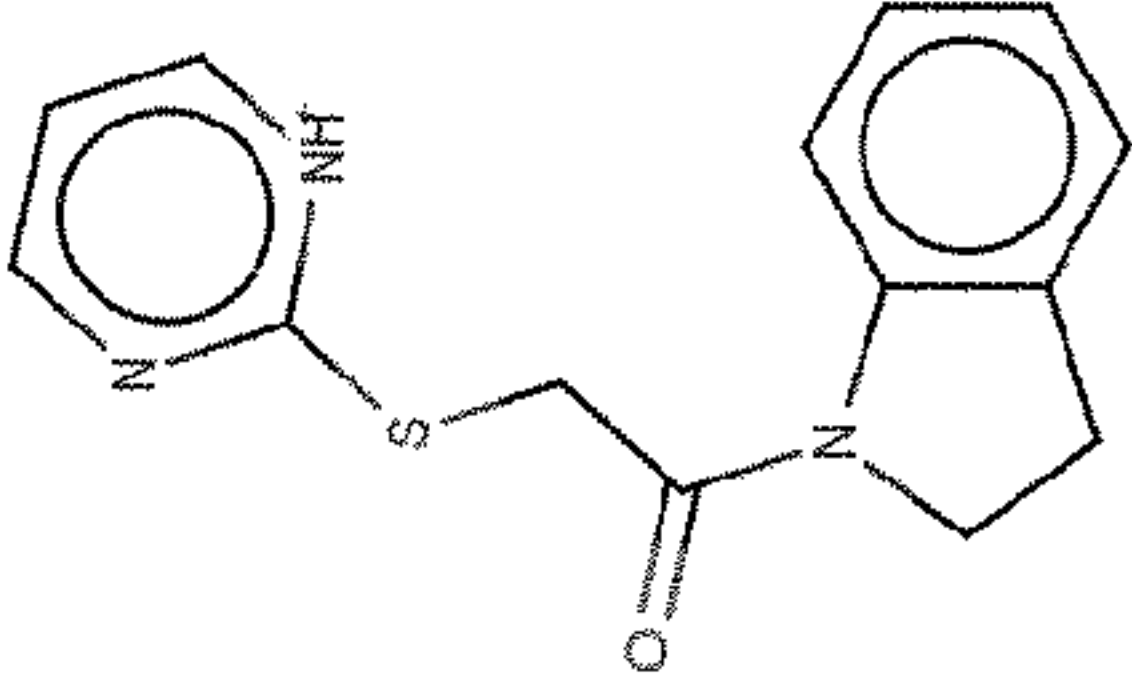
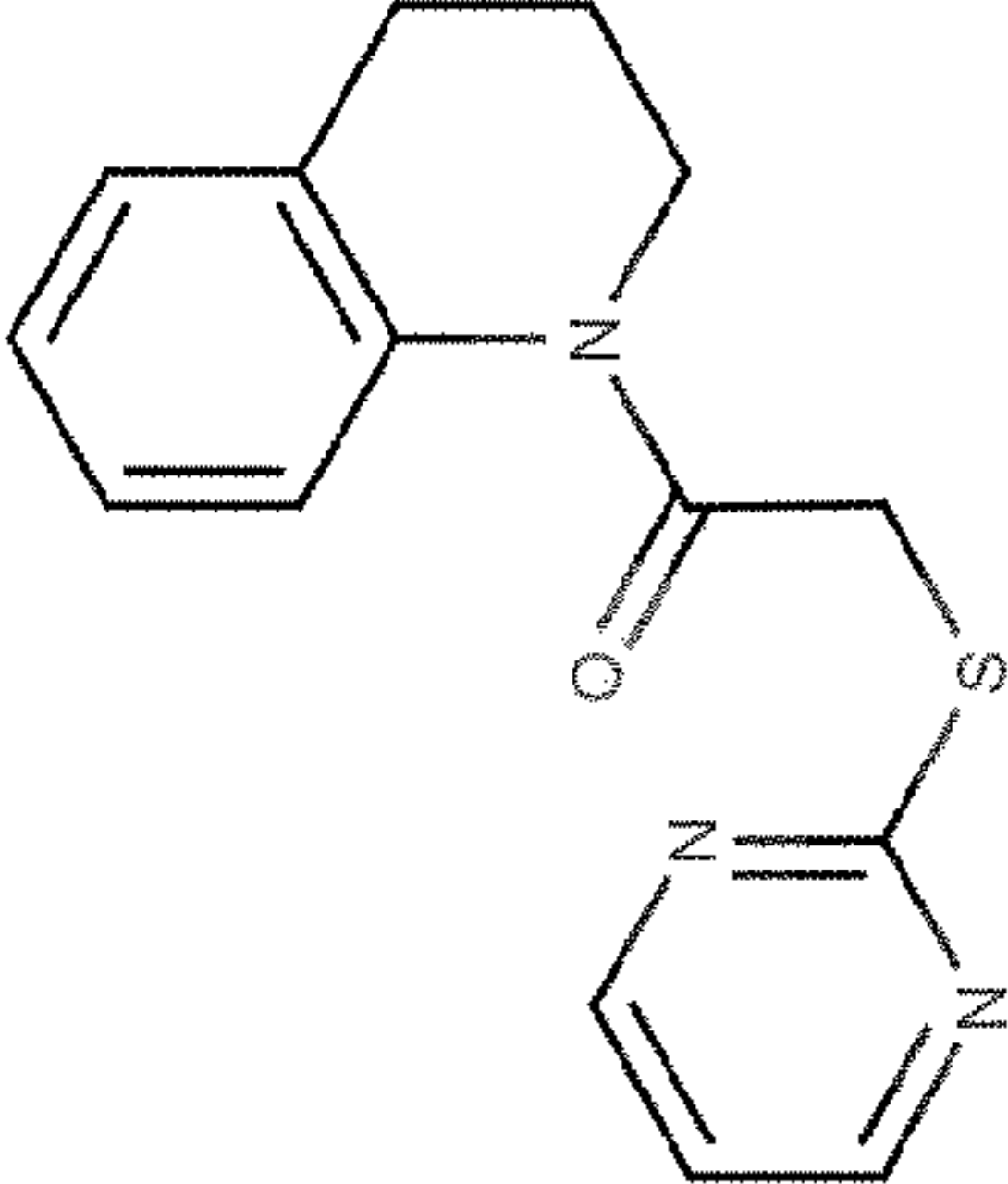
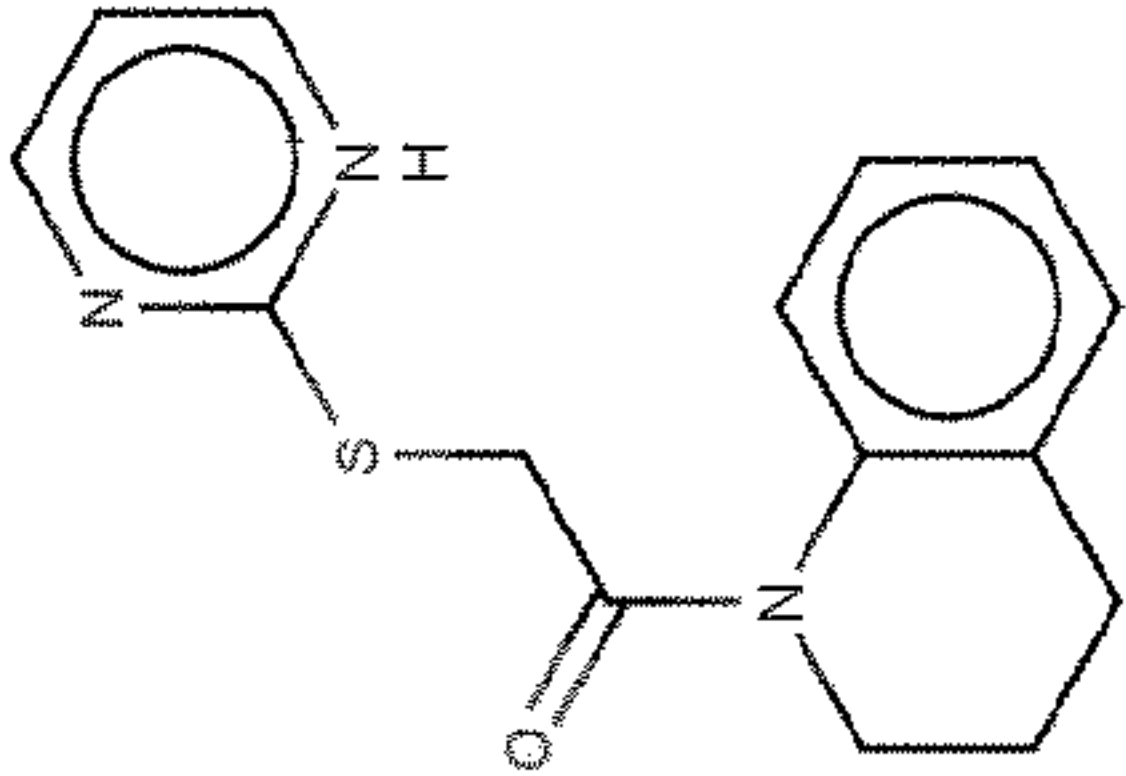
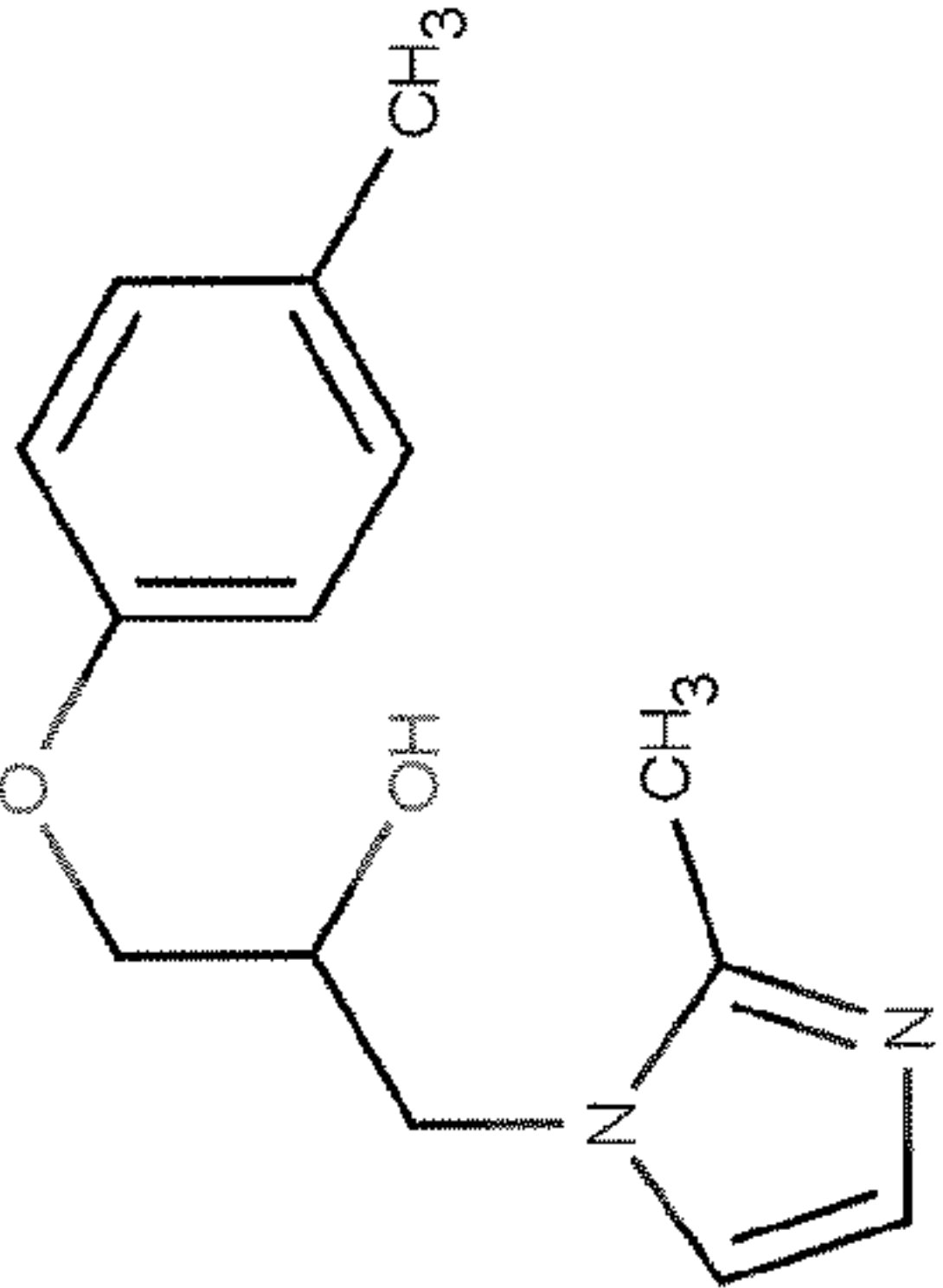
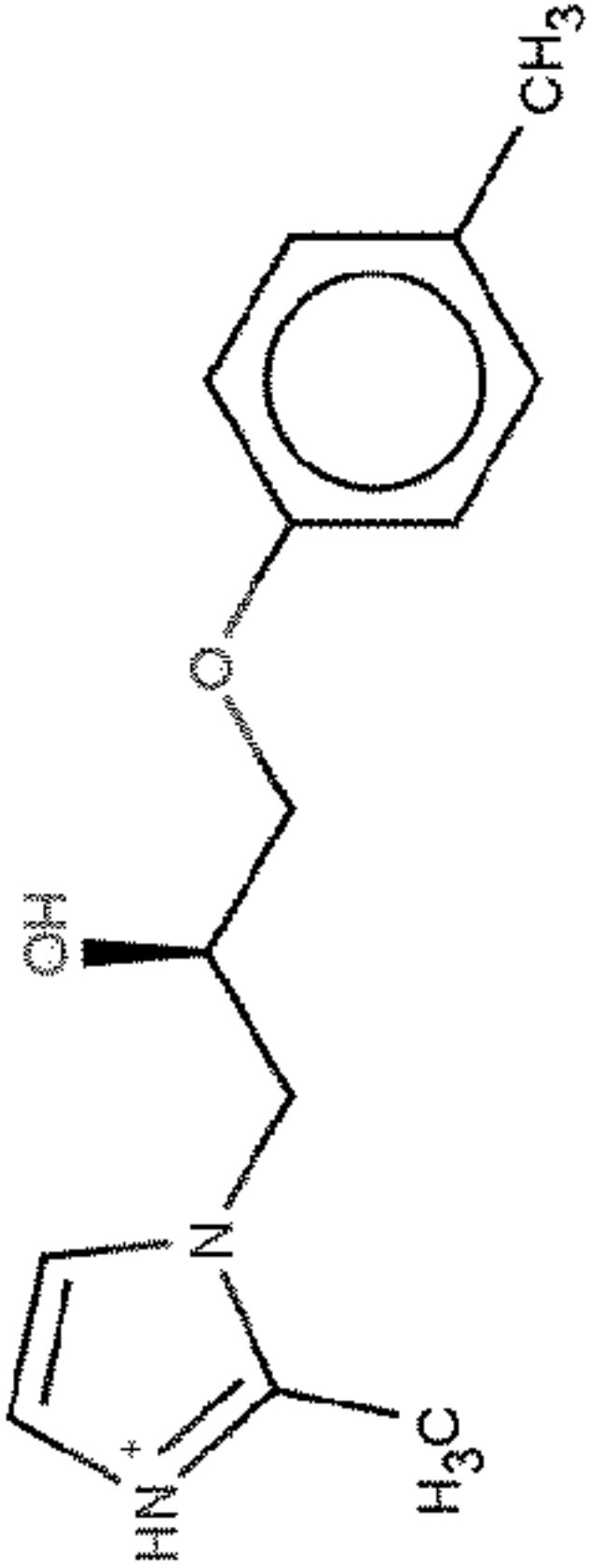
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 24		F0815-0092		F0815-0092_1_1 (1 of 7)
Compound 25		F0840-0033		F0840-0033_2 (1 of 4)
Compound 26		F0840-0034		F0840-0034_1_1 (1 of 4)

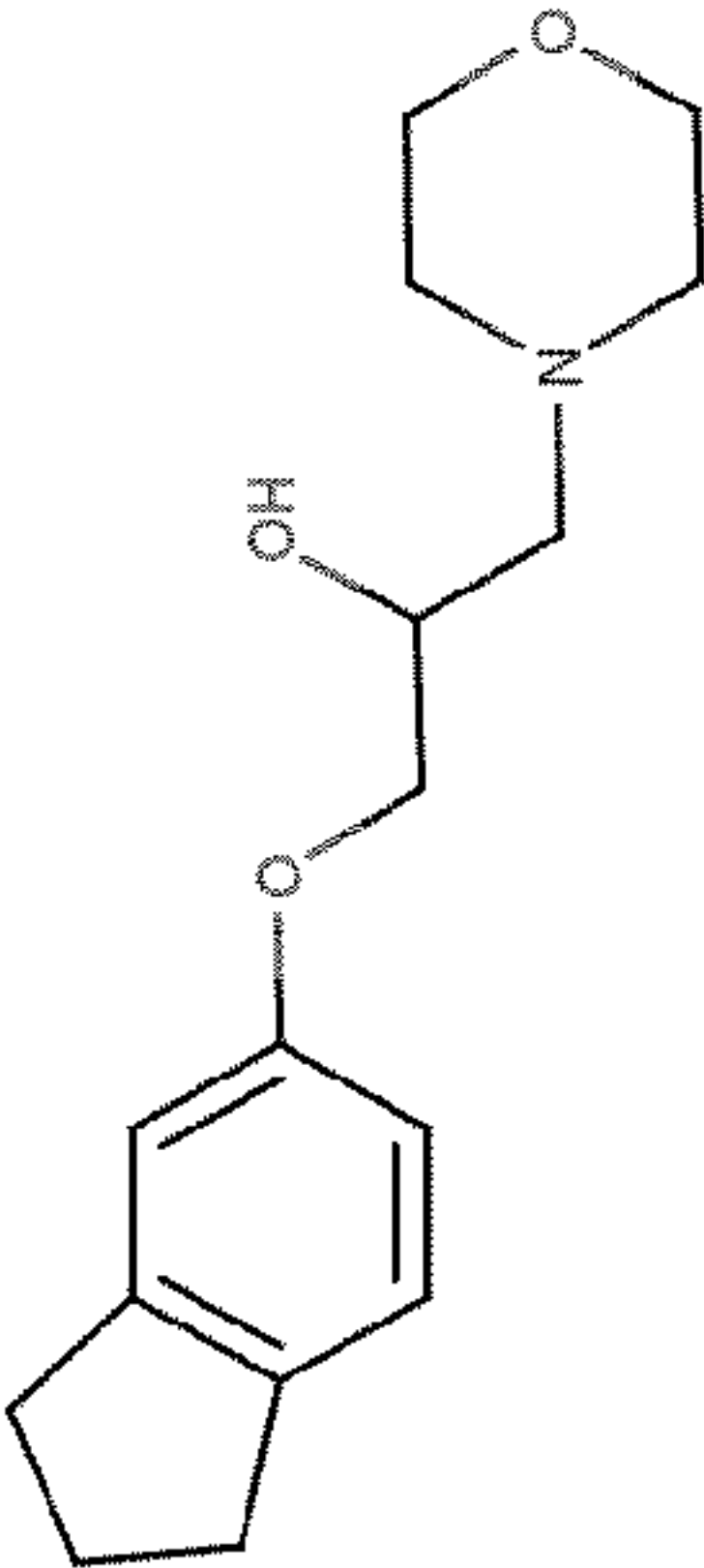
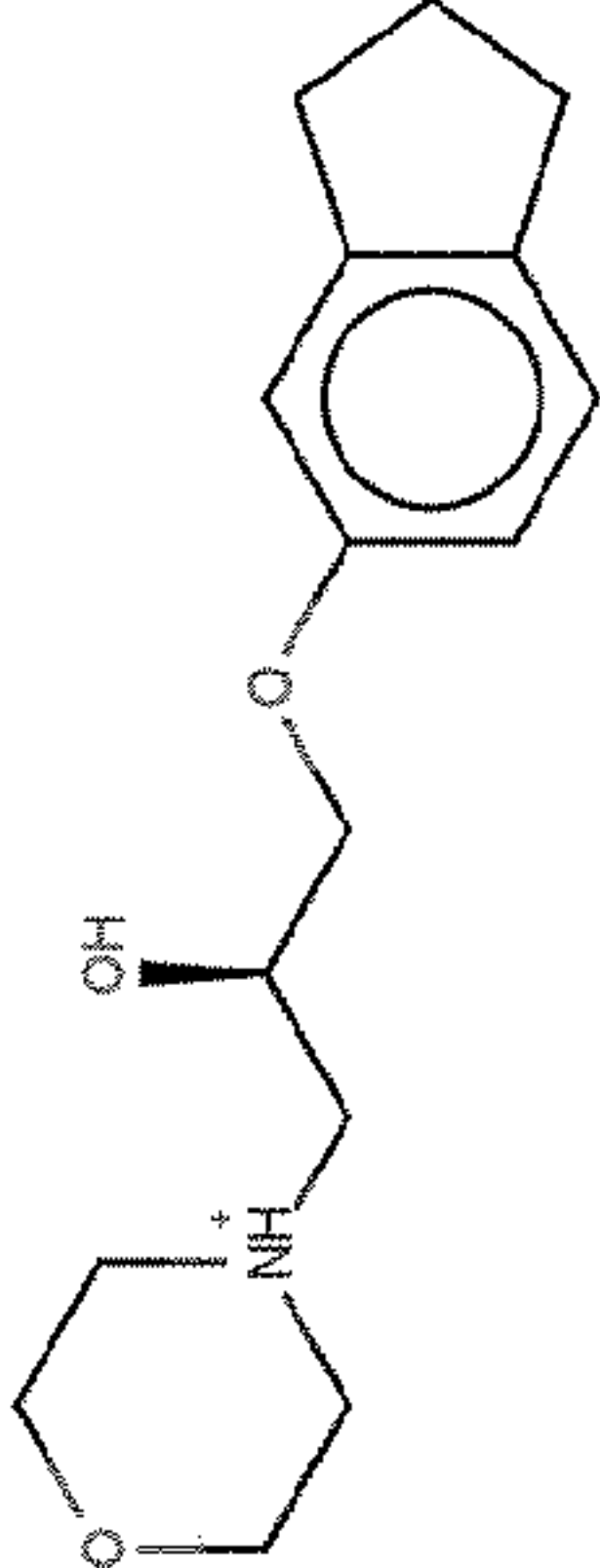
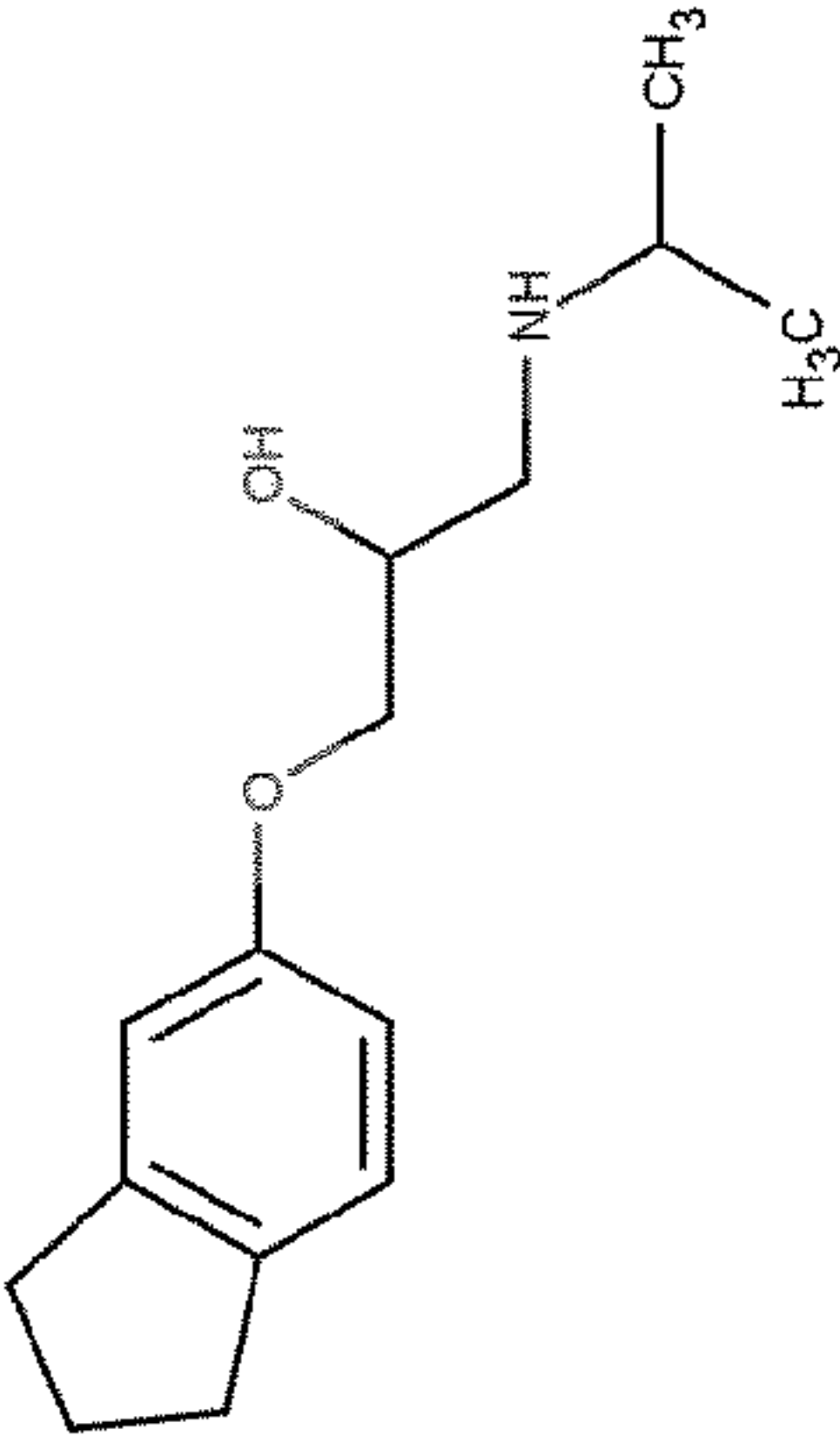
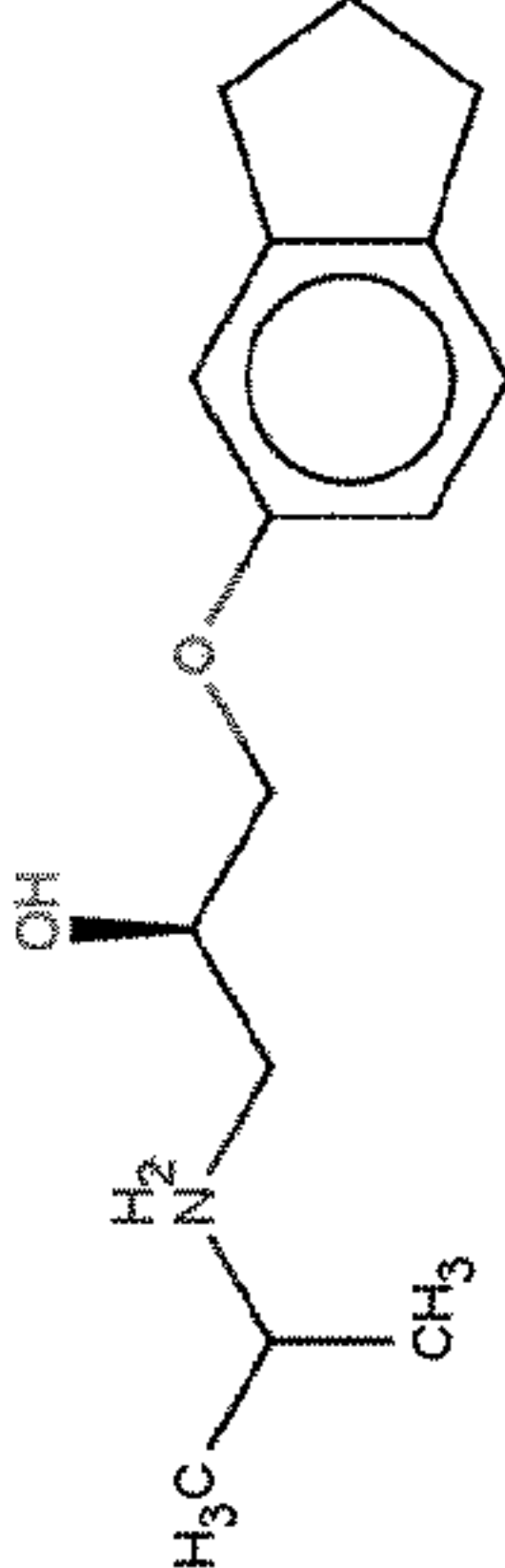
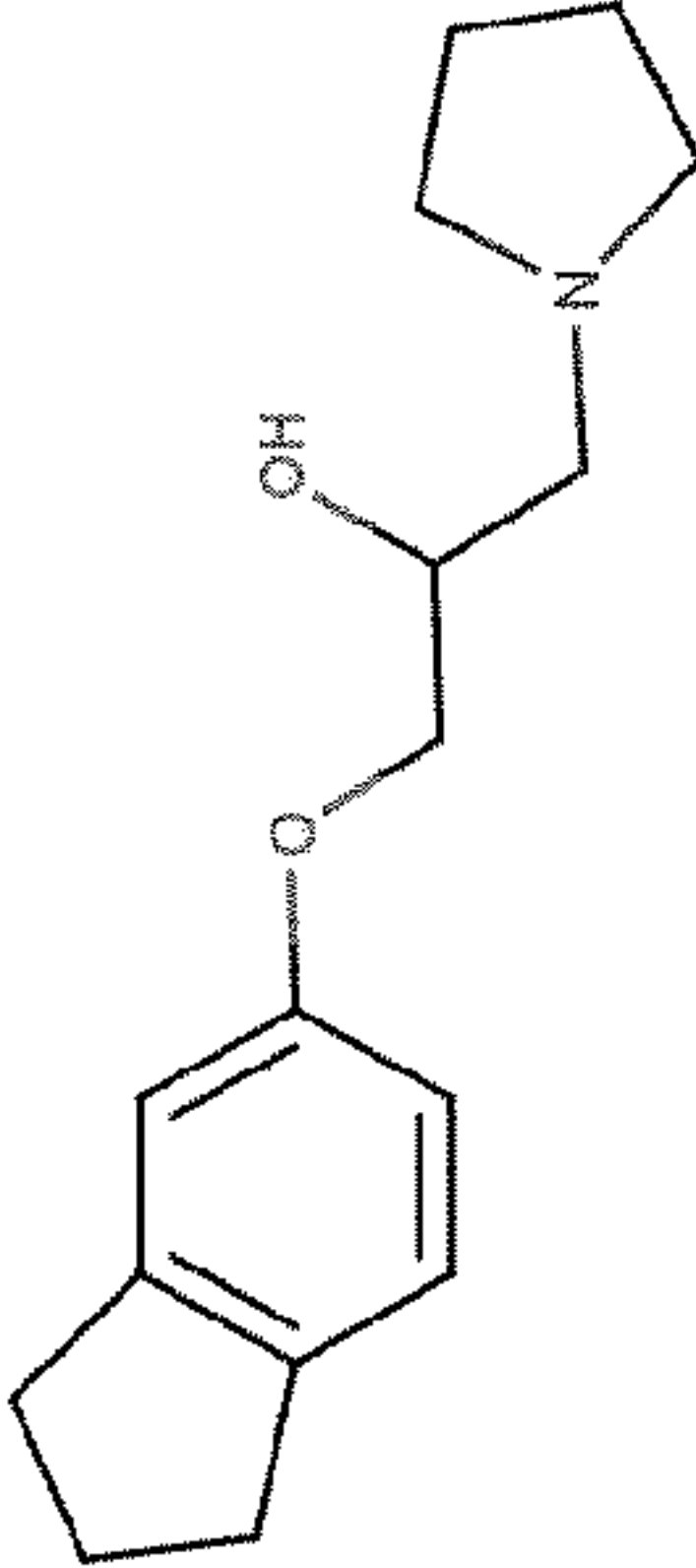
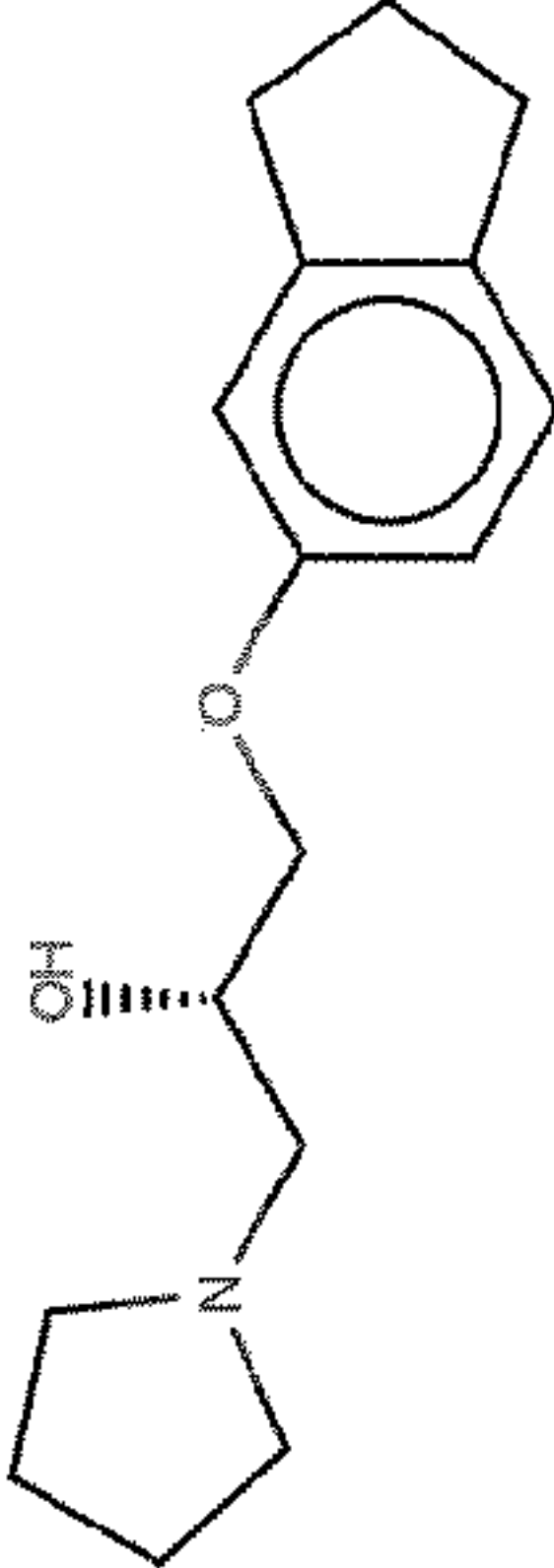
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 27		F0840-0058		F0840-0058_1 (1 of 4)
Compound 28		F0840-0059		F0840-0059_1 (1 of 8)
Compound 29		F0840-0204		F0840-0204_1_2 (1 of 4)

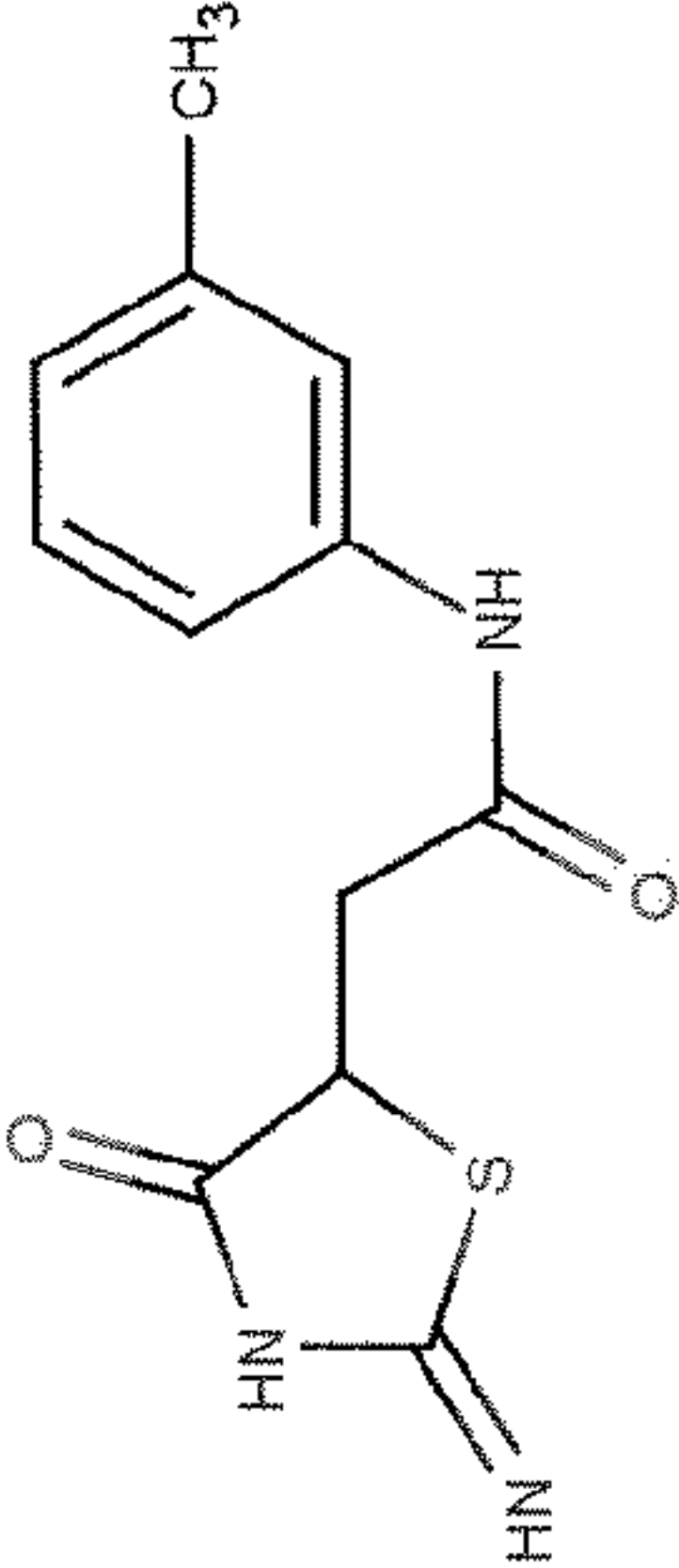
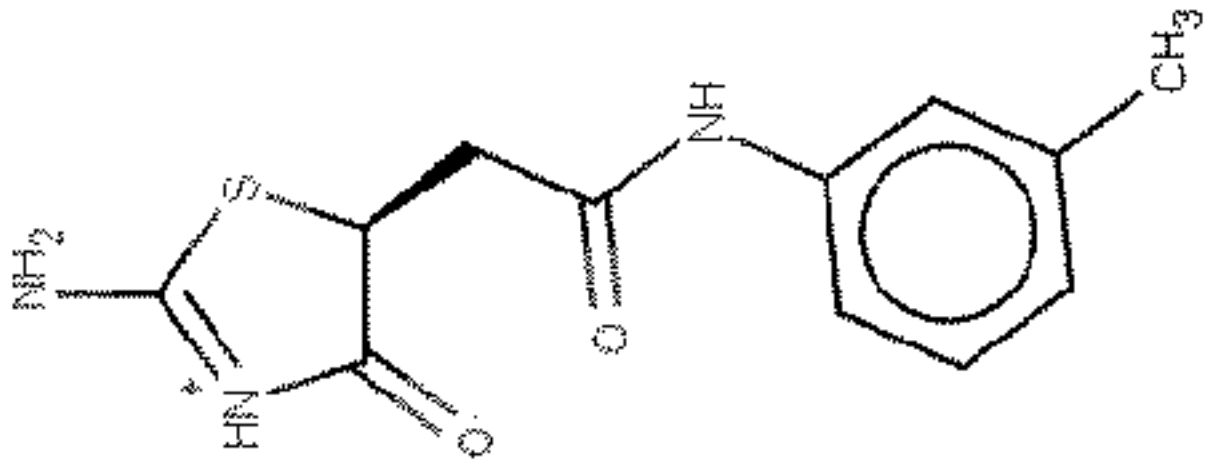
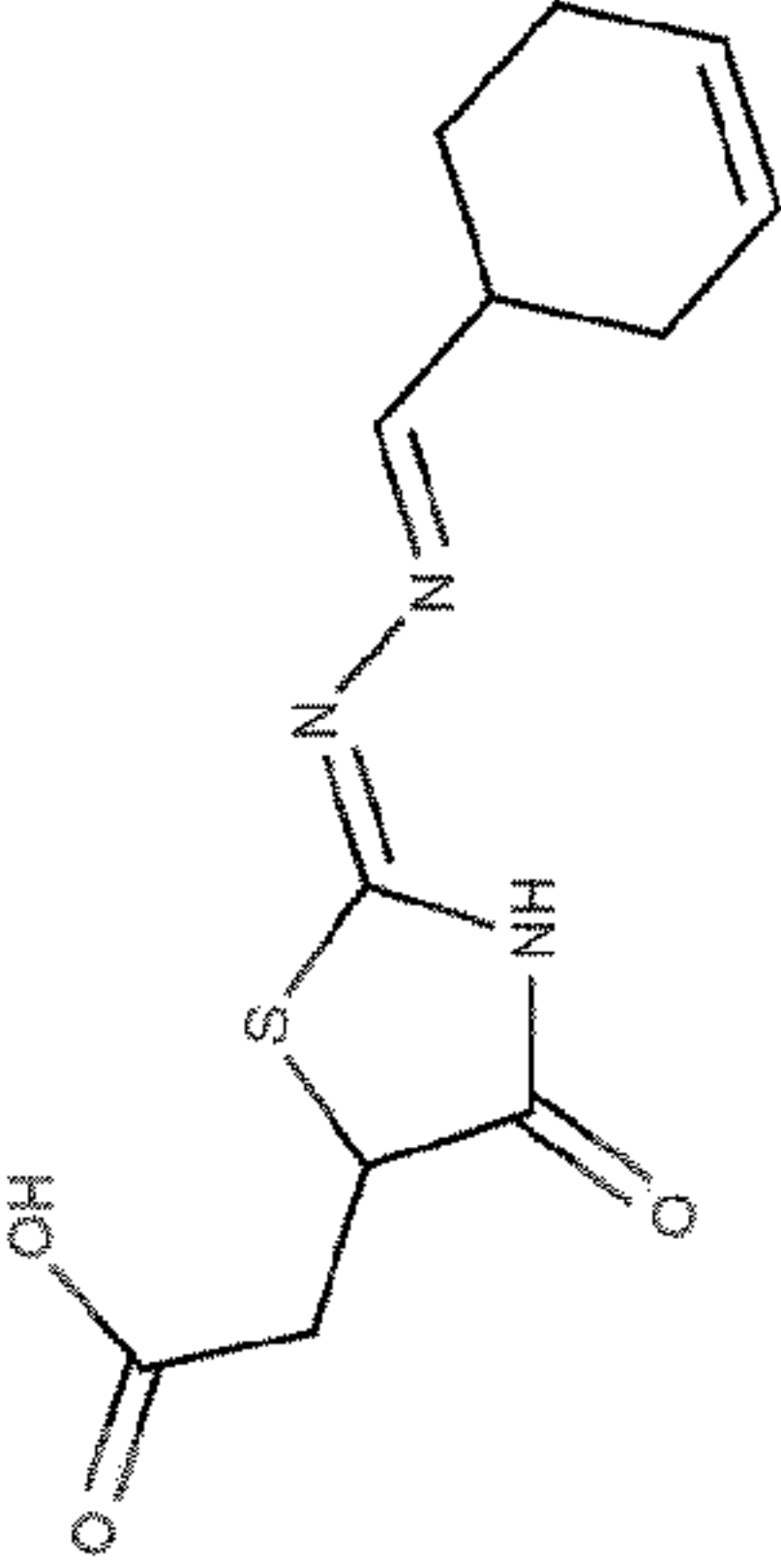
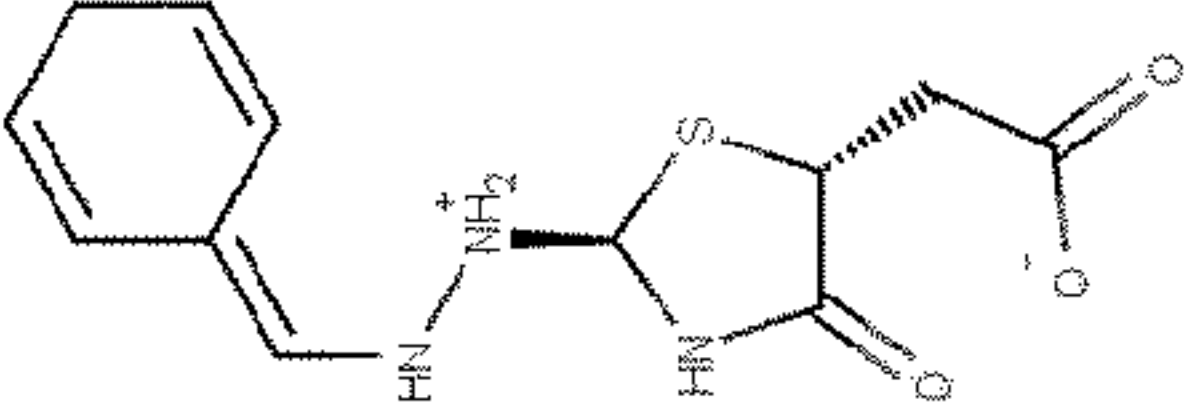
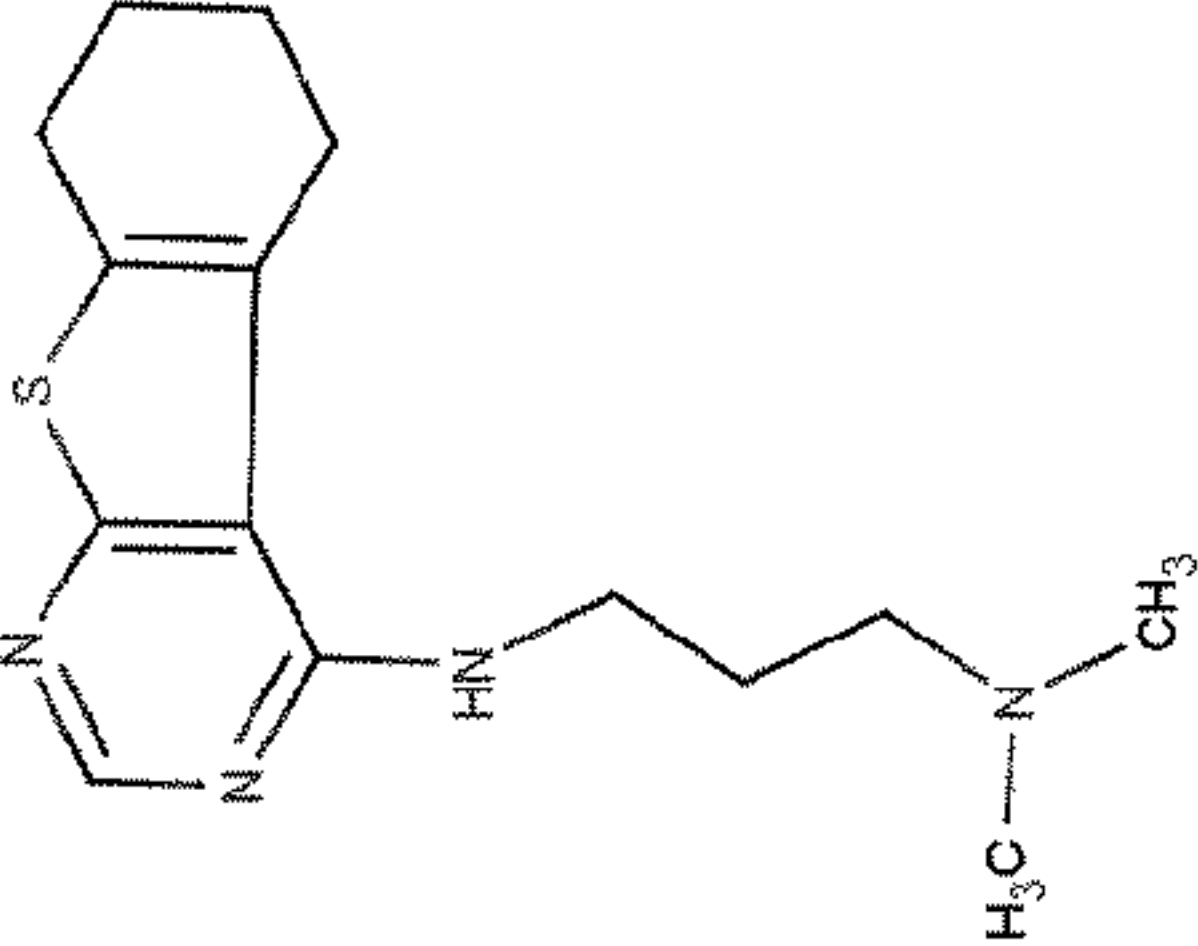
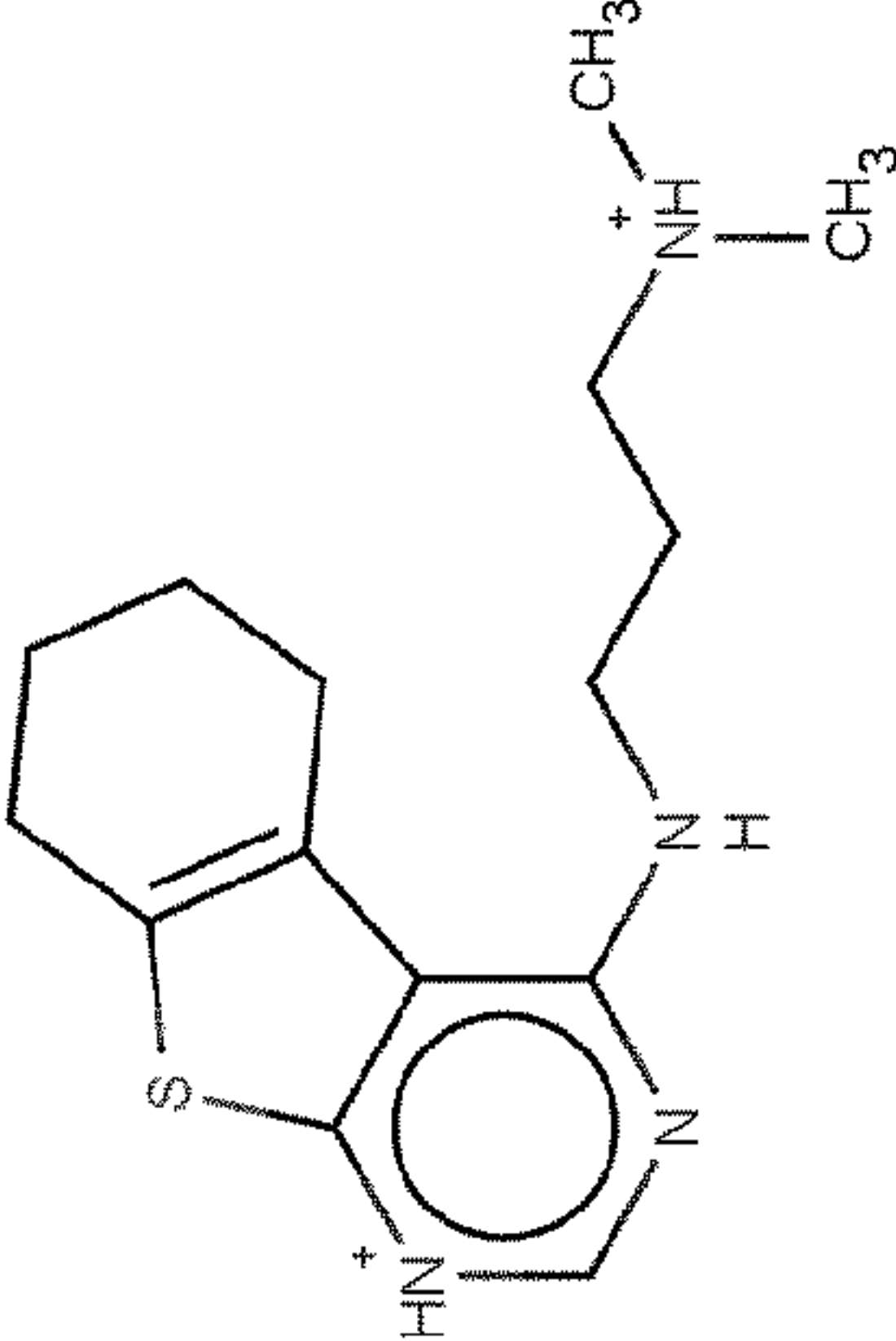
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 30		F0840-0206		F0840-0206_2 (1 of 4)
Compound 31		F0840-0214		F0840-0214_2 (1 of 4)
Compound 32		F0896-0201		F0896-0201_1_2 (1 of 4)

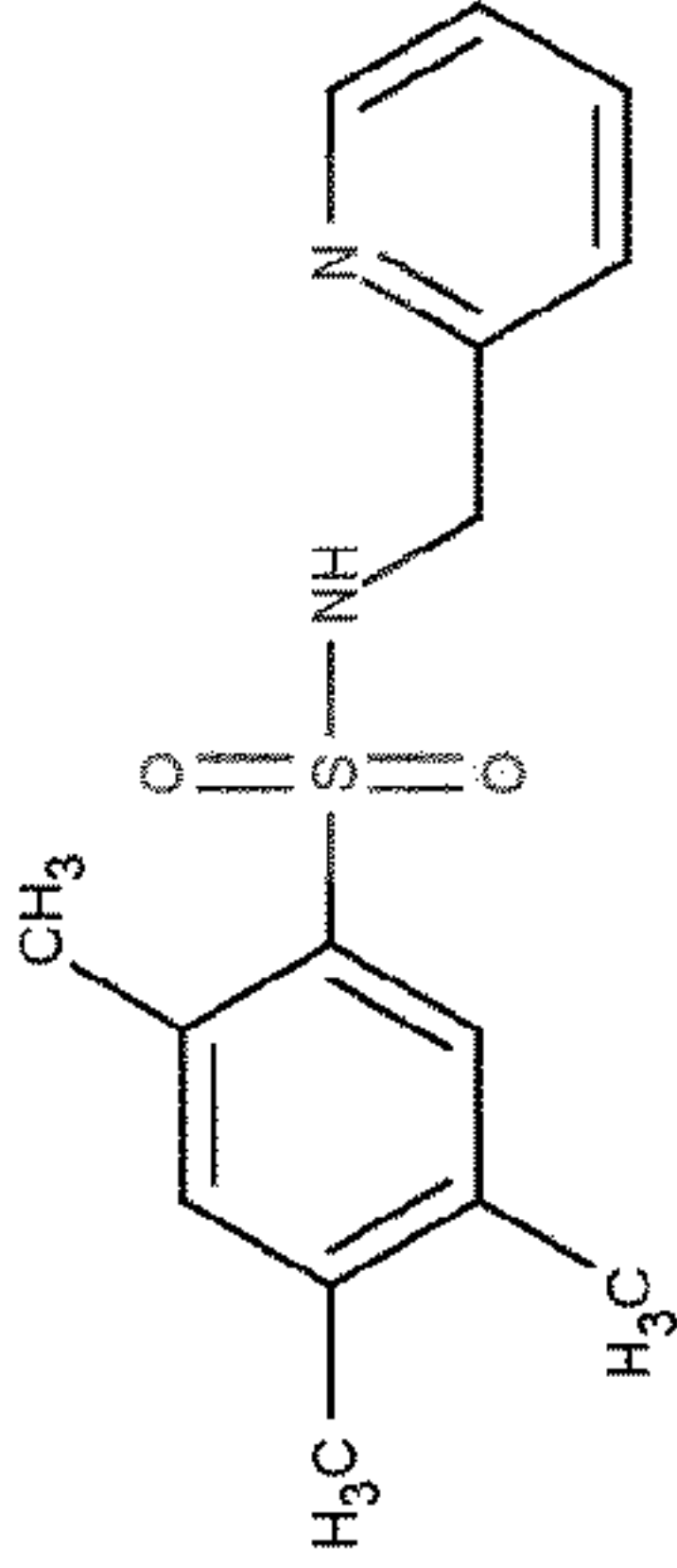
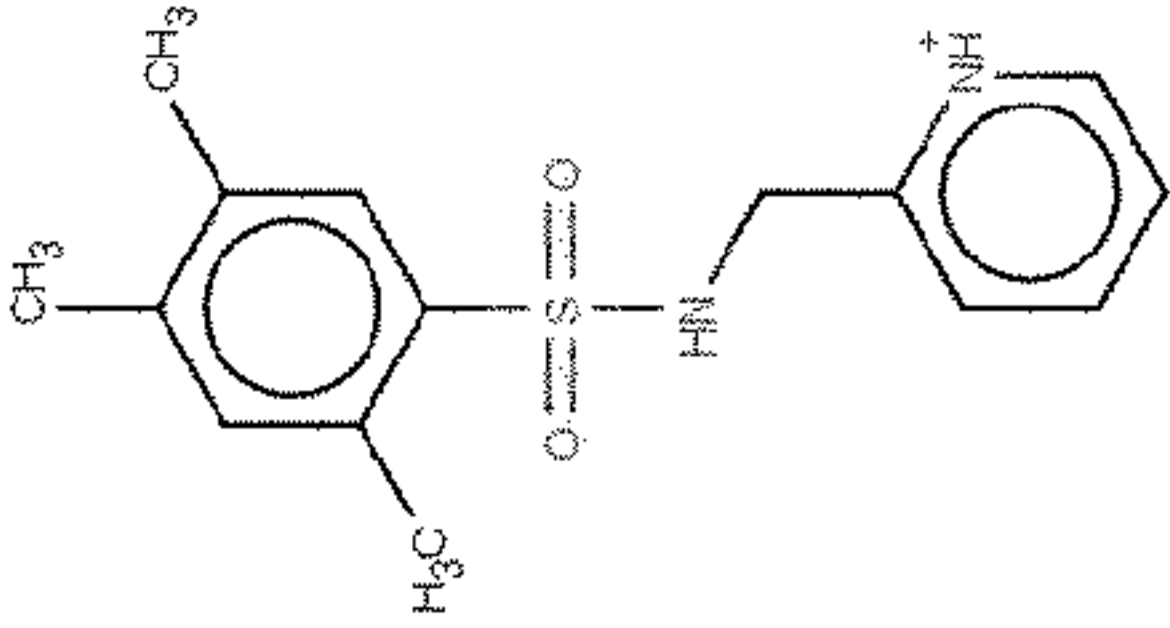
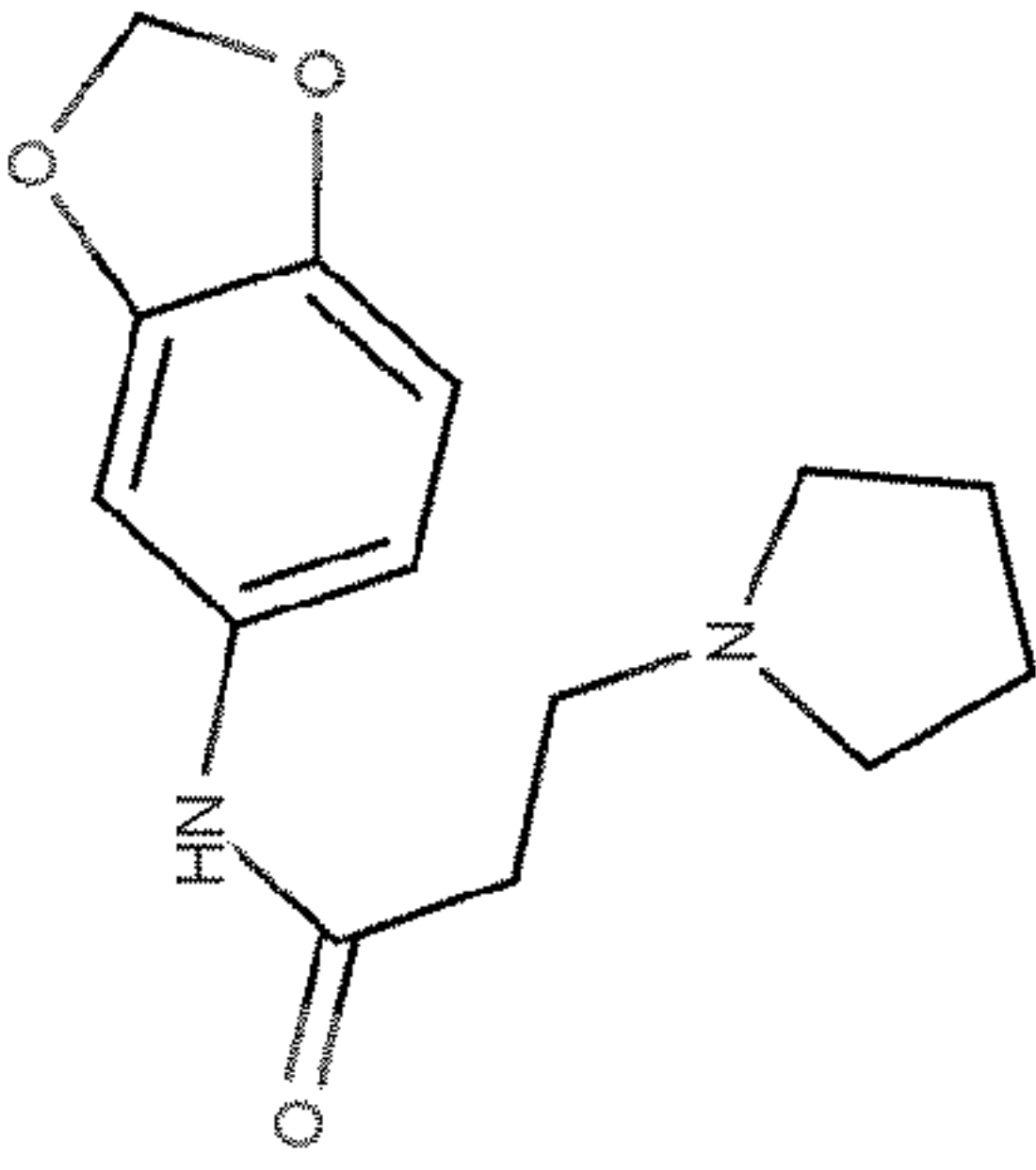
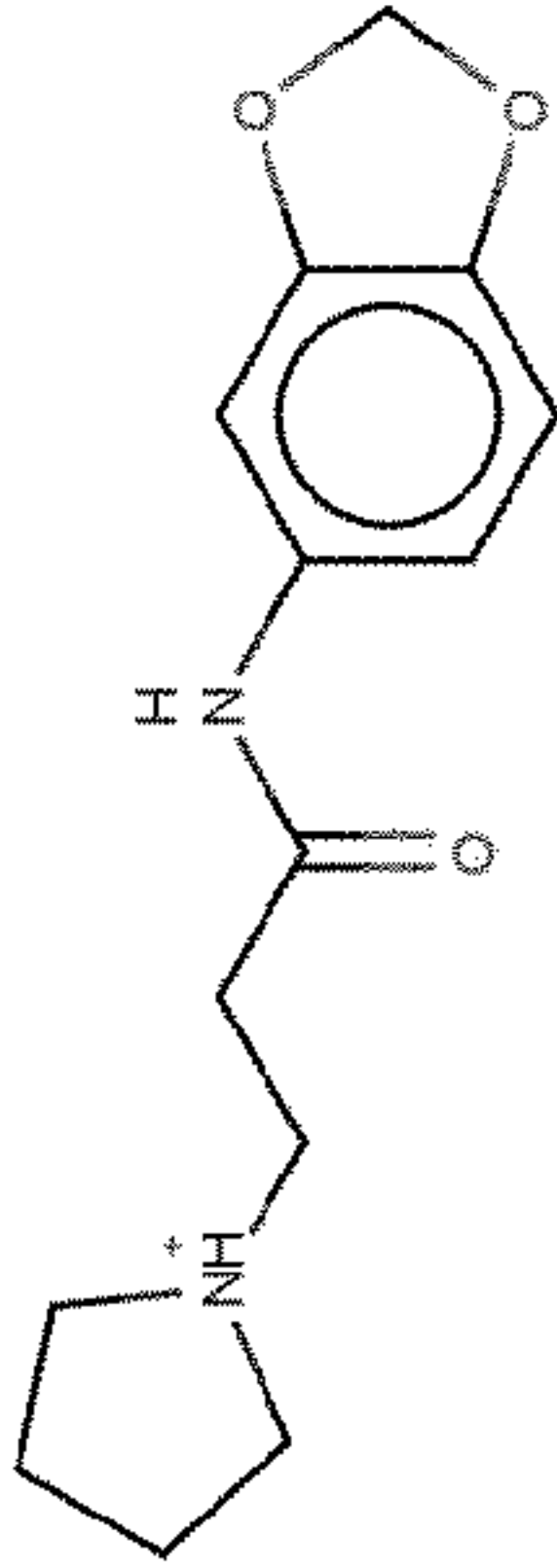
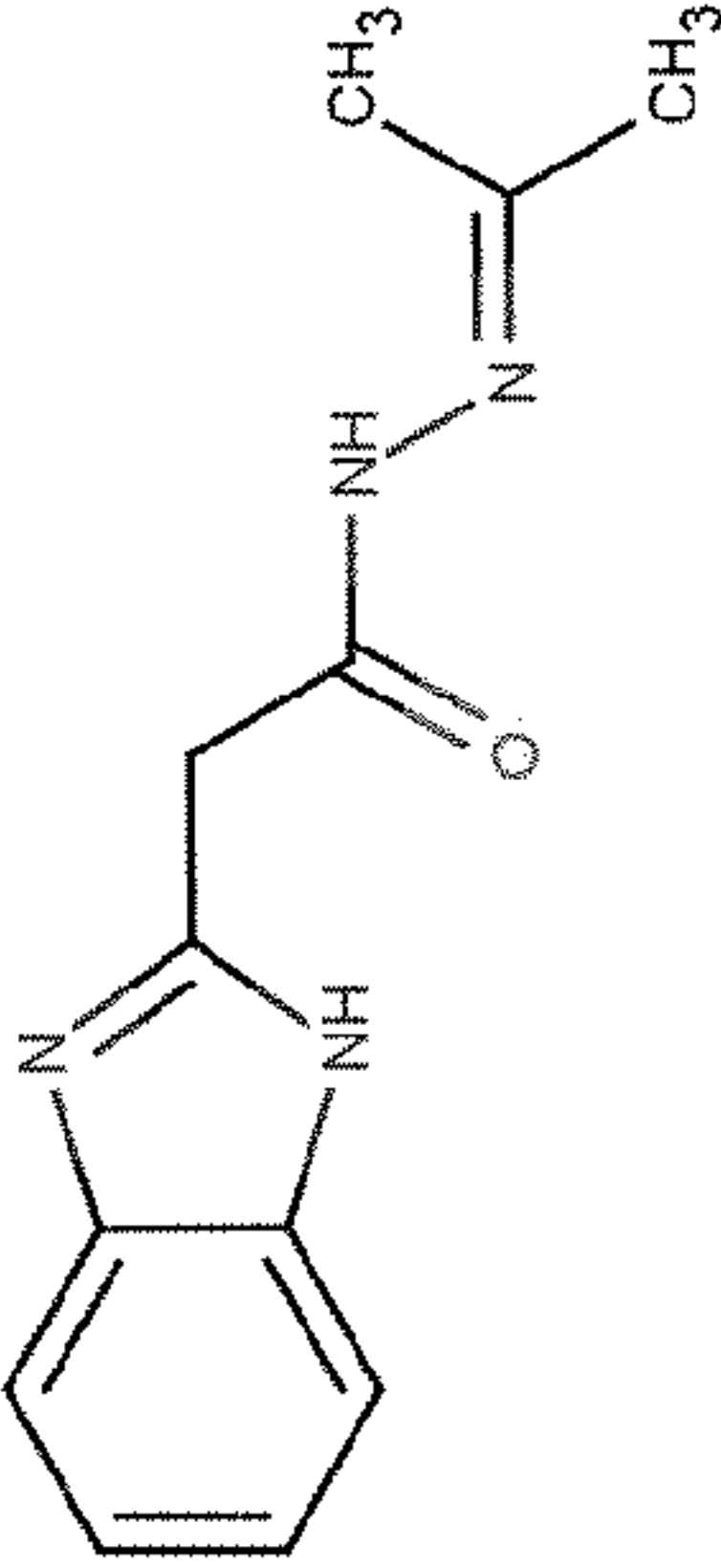
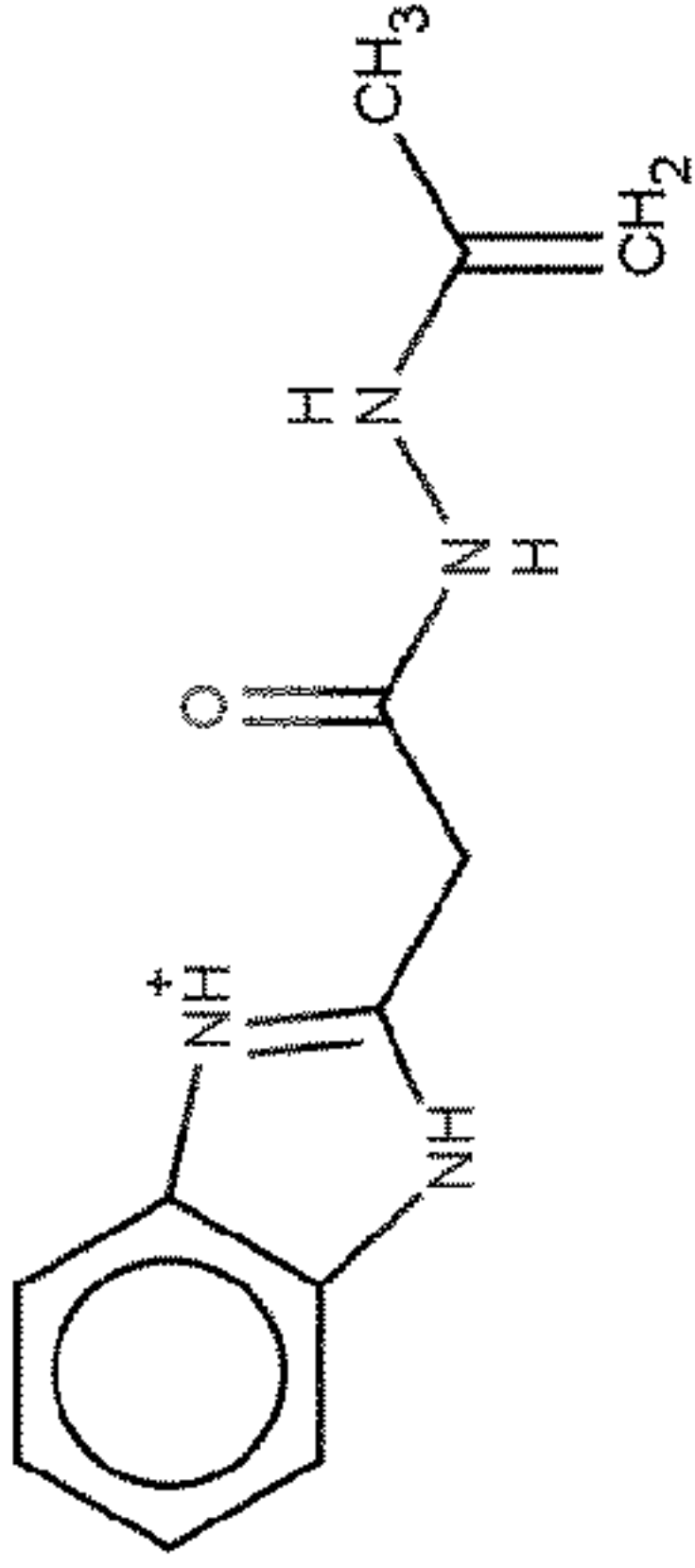
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 33		F0910-6535		F0910-6535_23_1 (1 of 124)
Compound 34		F0915-2932		F0915-2932_3_1 (1 of 8)
Compound 35		F0915-2972		F0915-2972_3_1 (1 of 5)

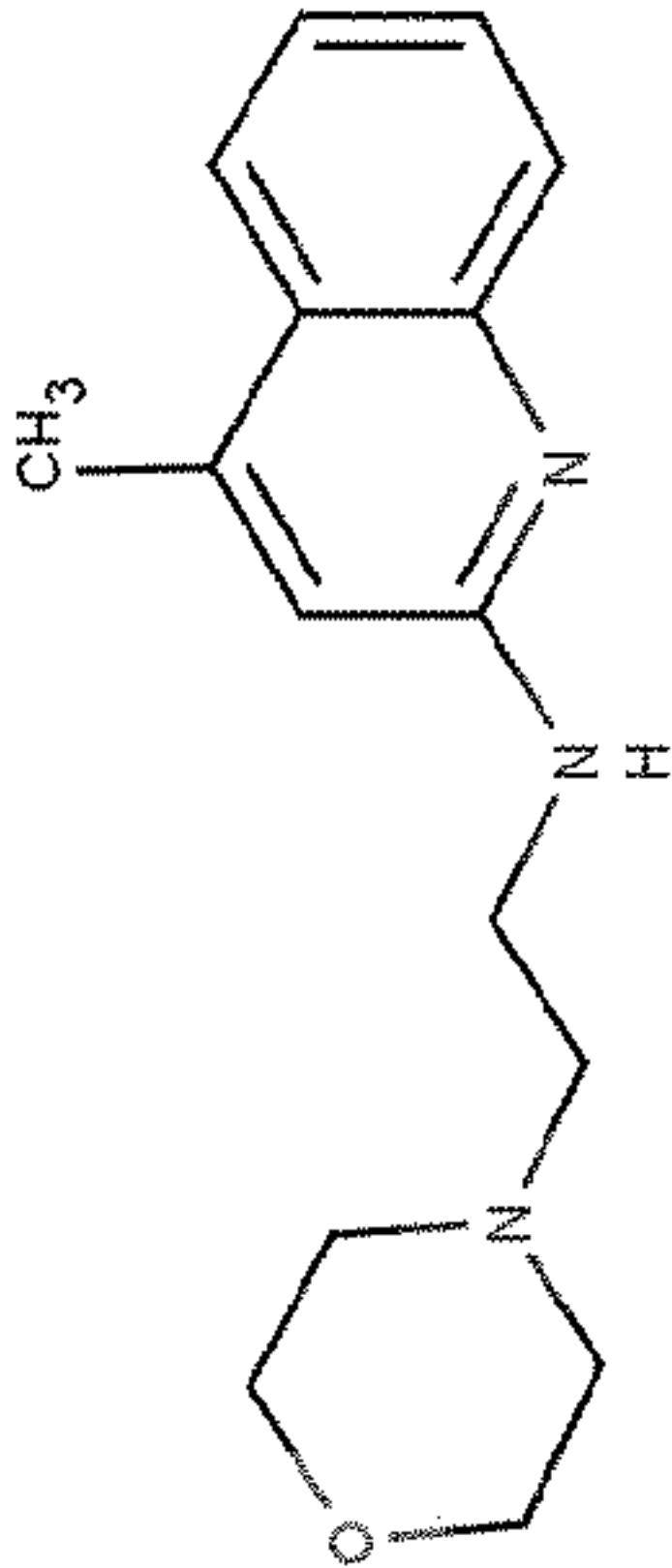
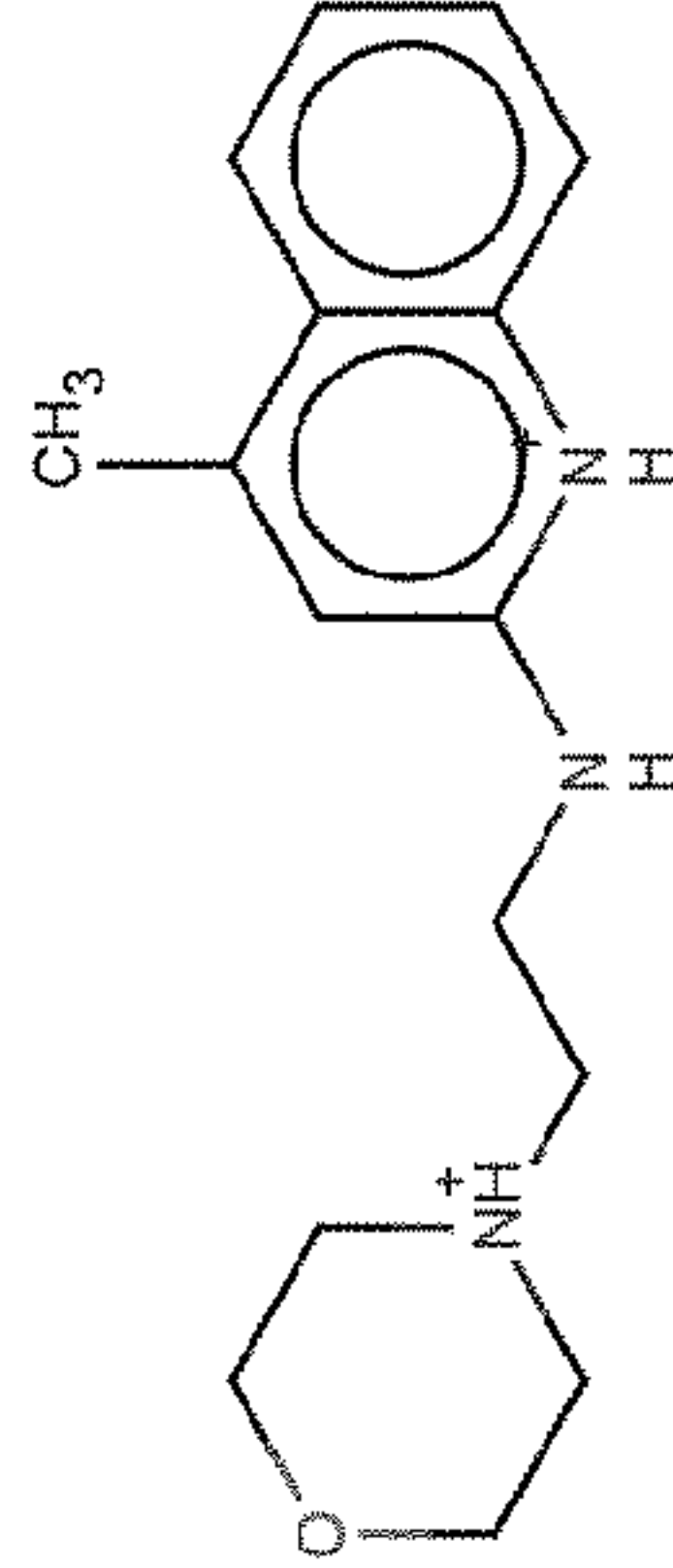
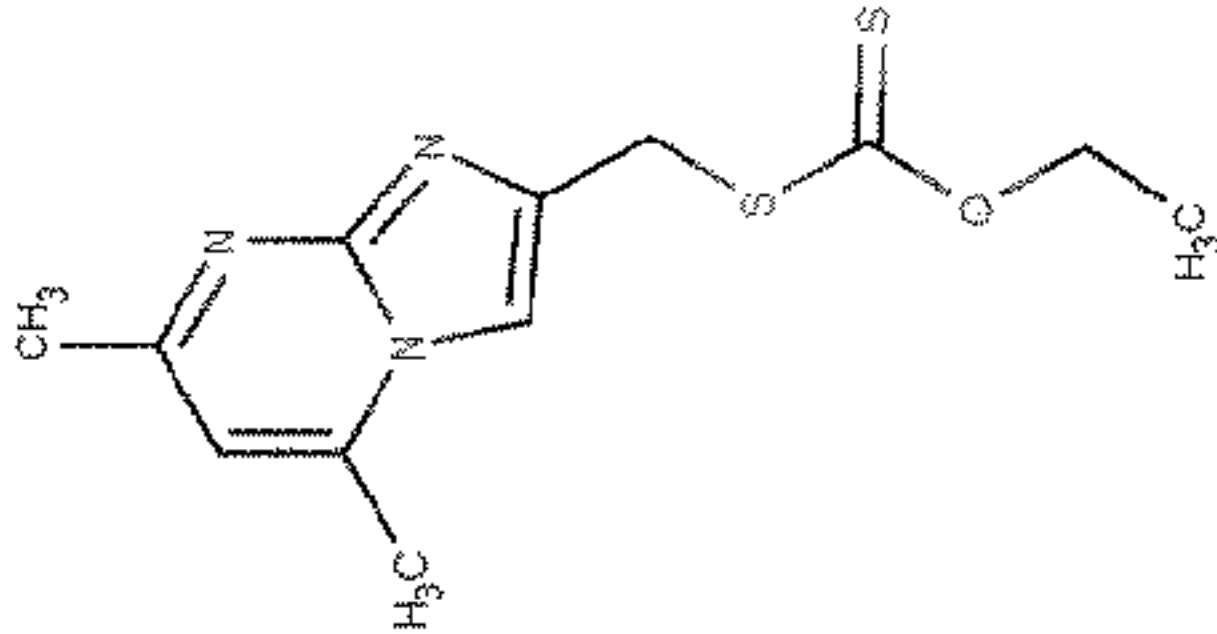
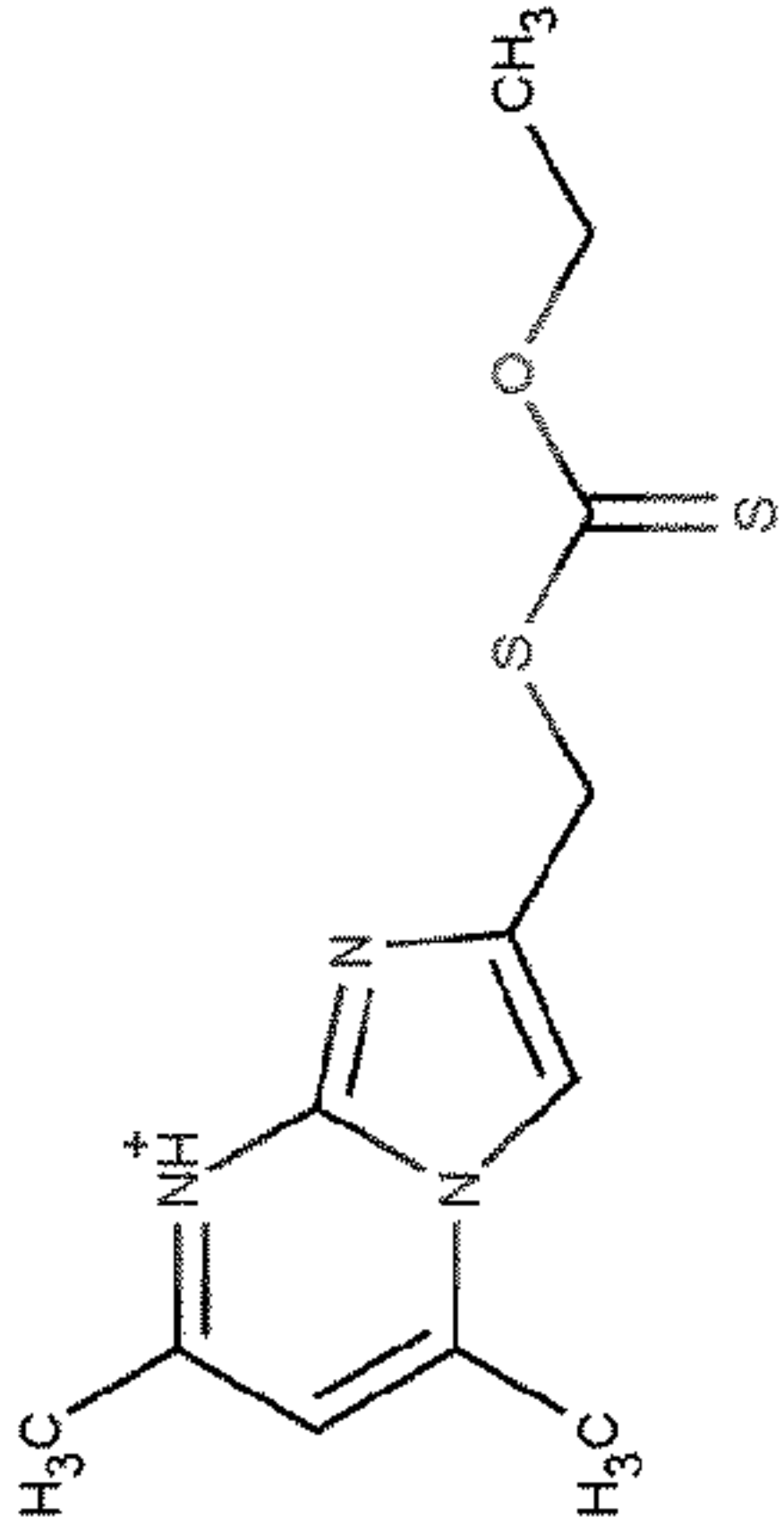
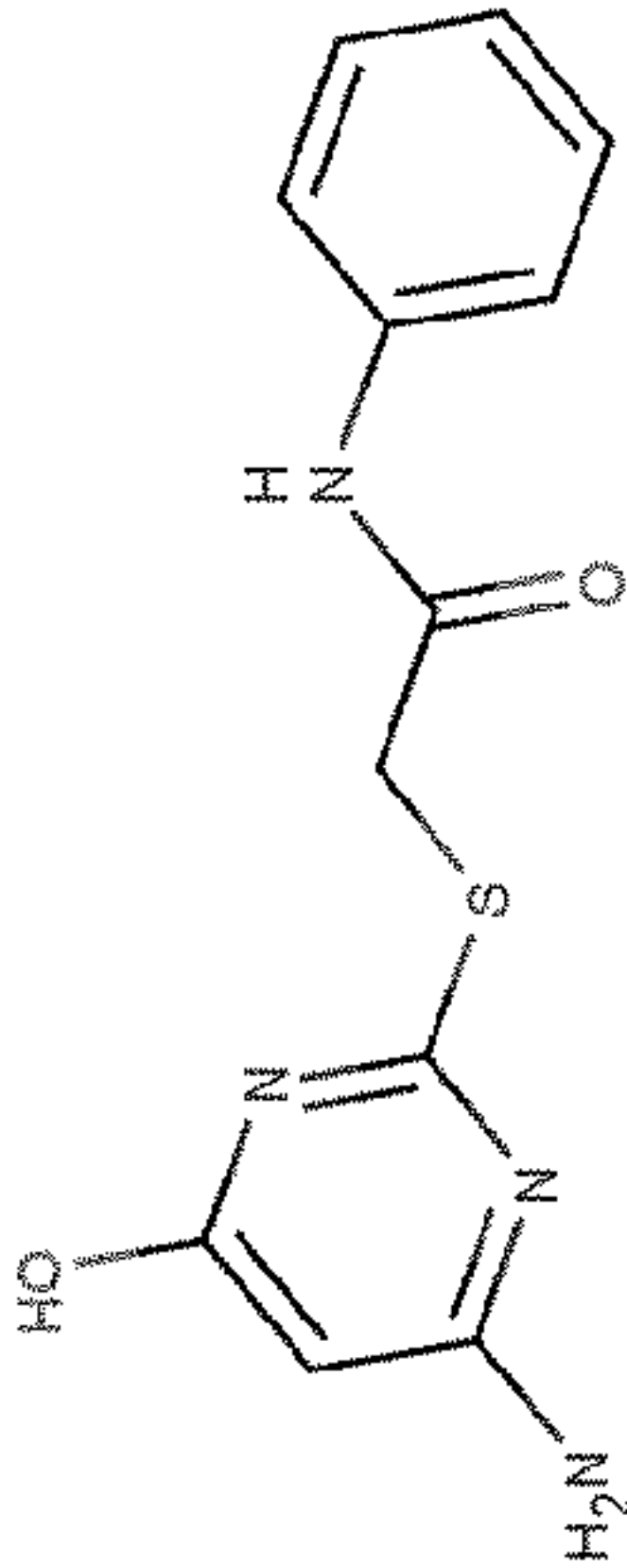
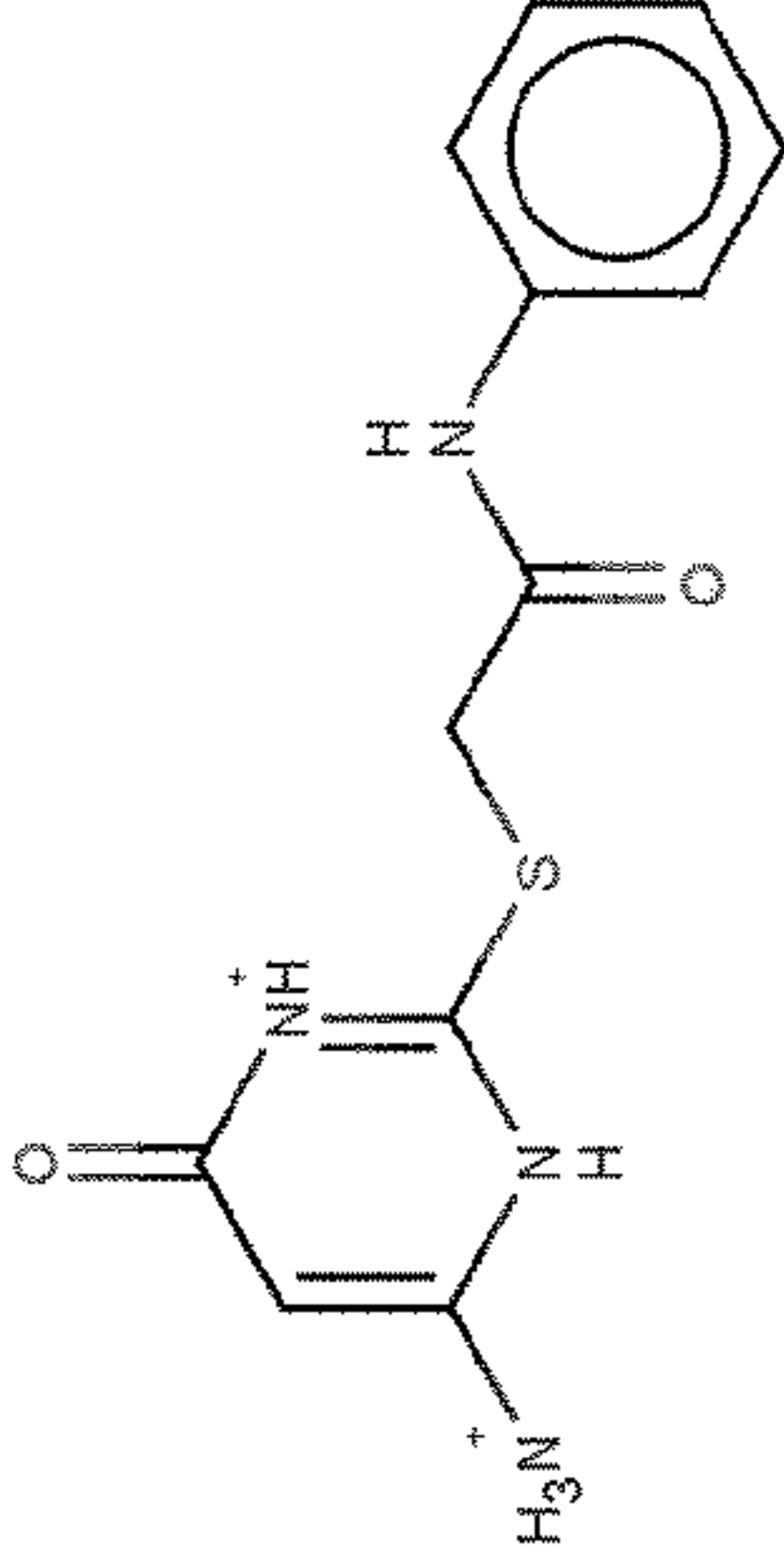
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 36		F0918-1472		F0918-1472_8_1 (1 of 10)
Compound 37		F0919-7794		F0919-7794_104_3 (1 of 270)
Compound 38		F1001-0002		F1001-0002_5_1 (1 of 12)

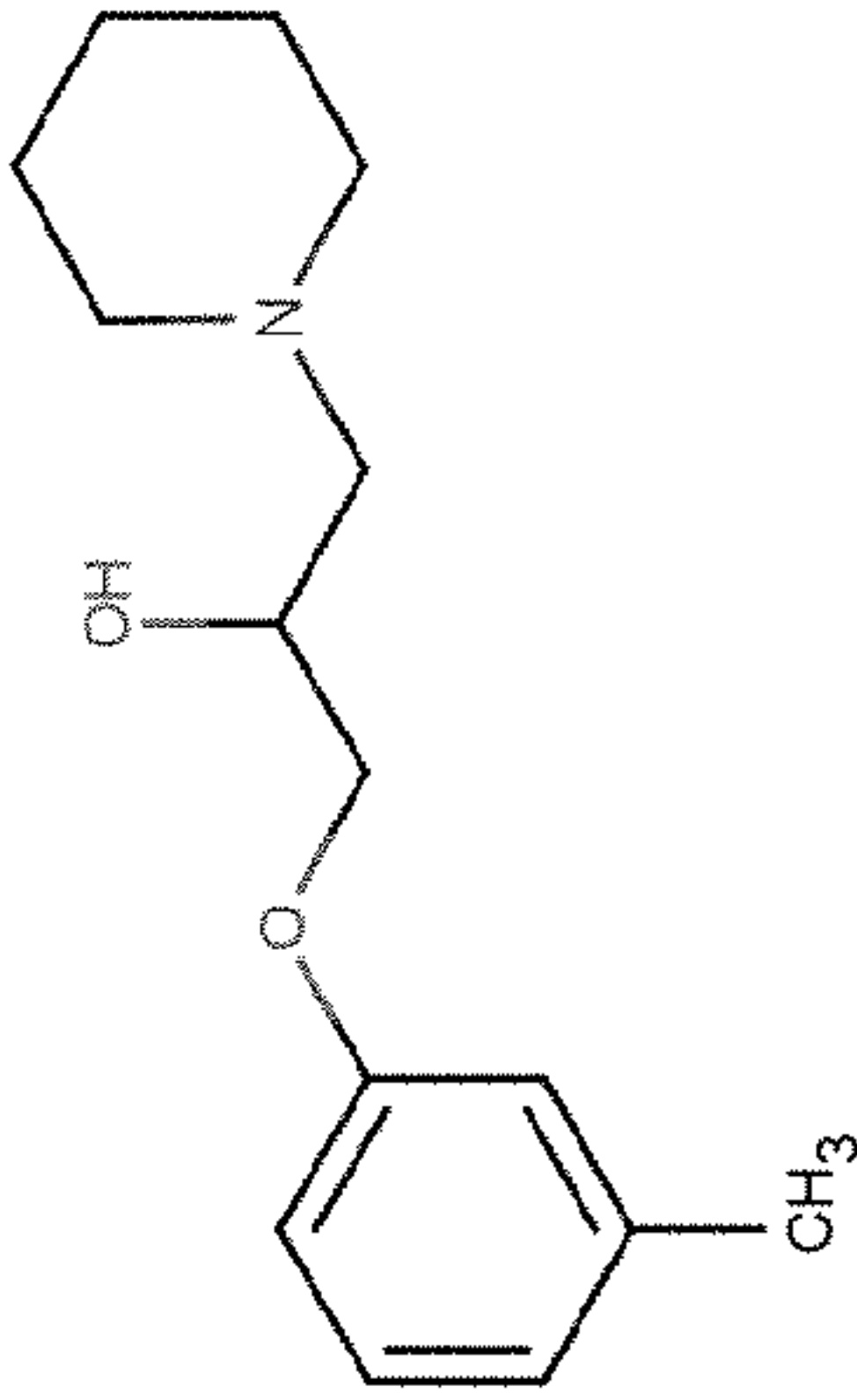
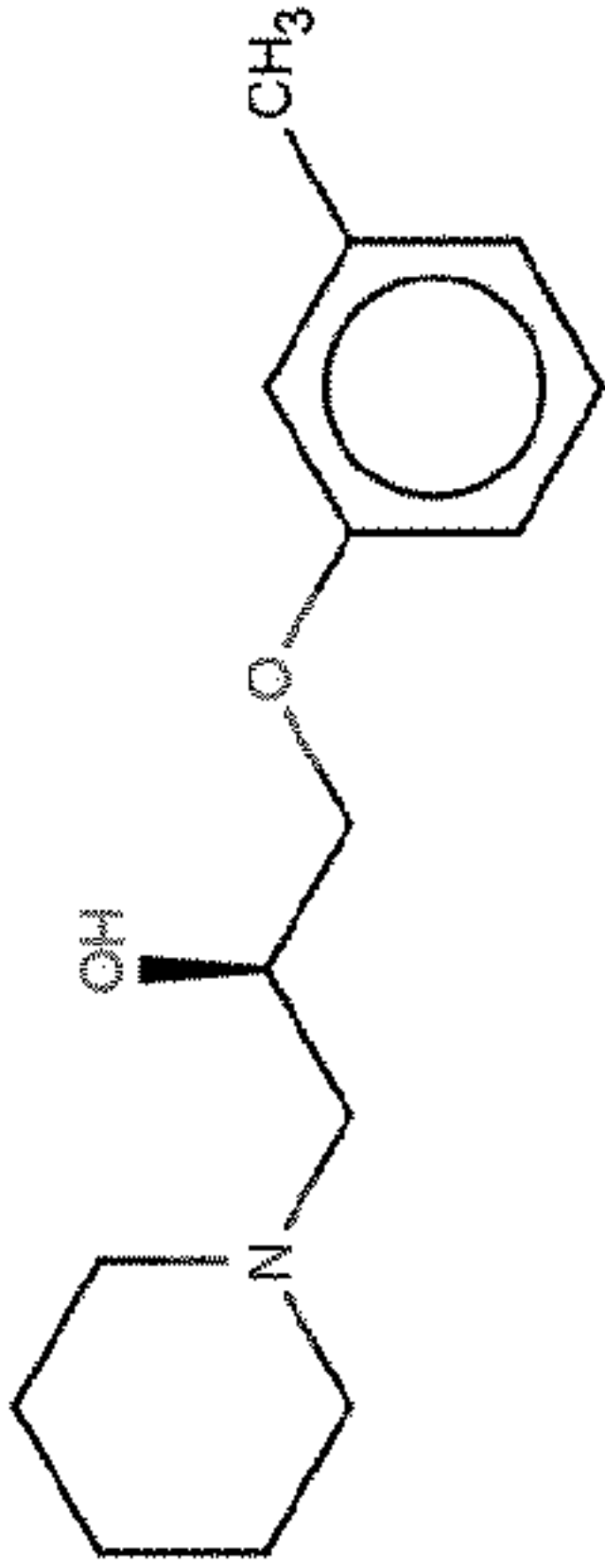
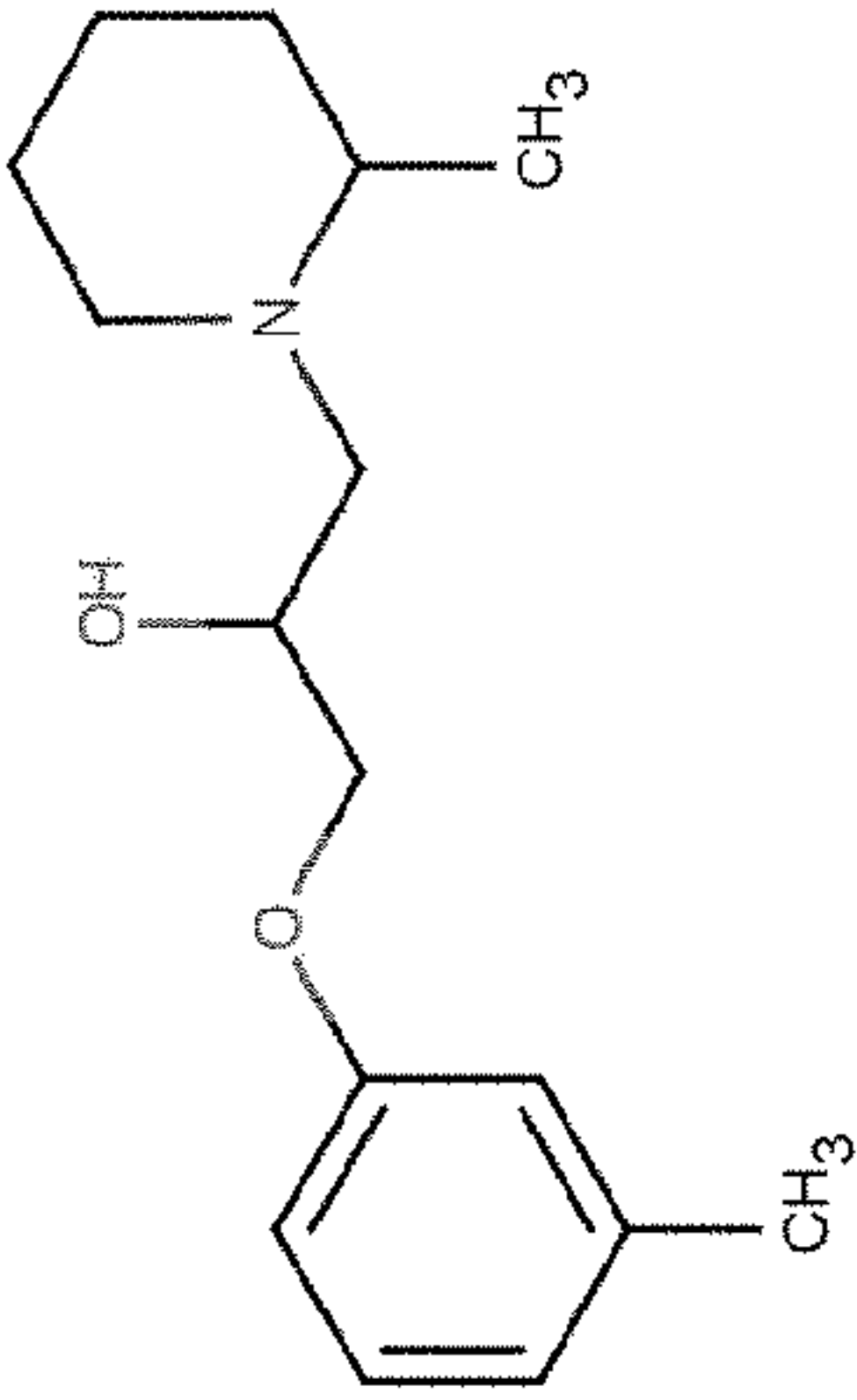
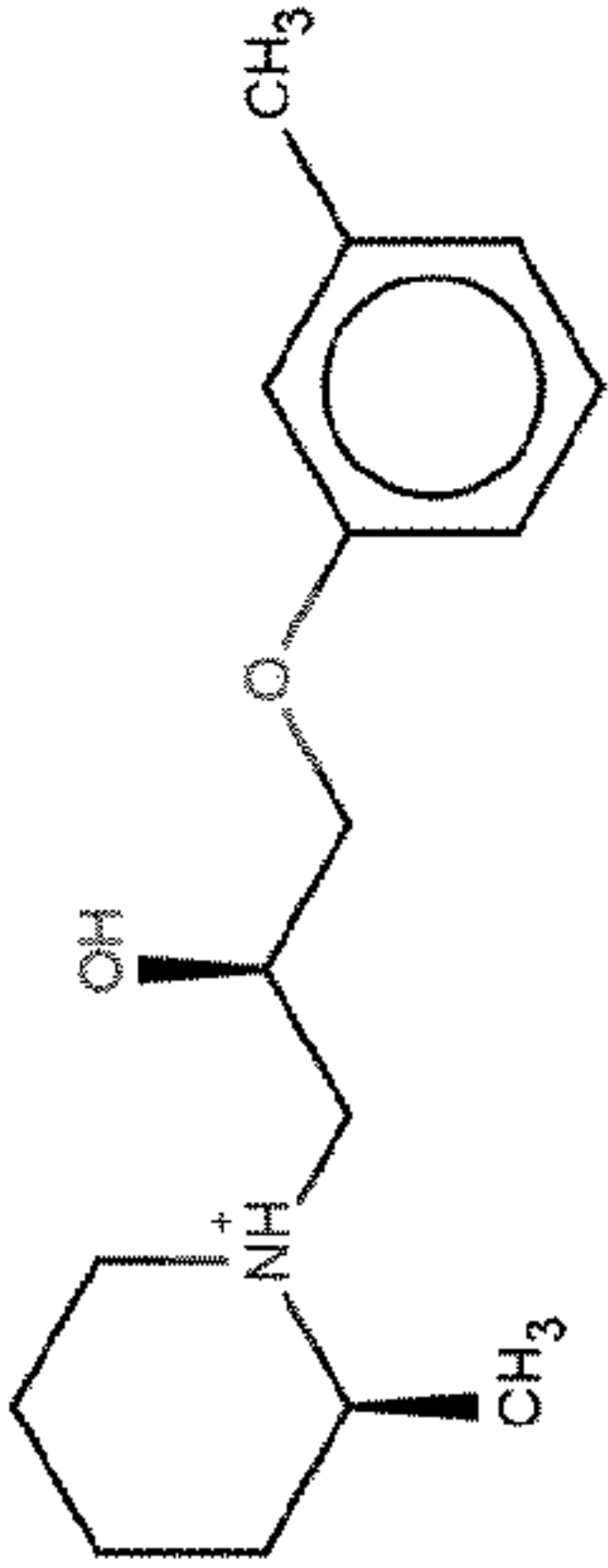
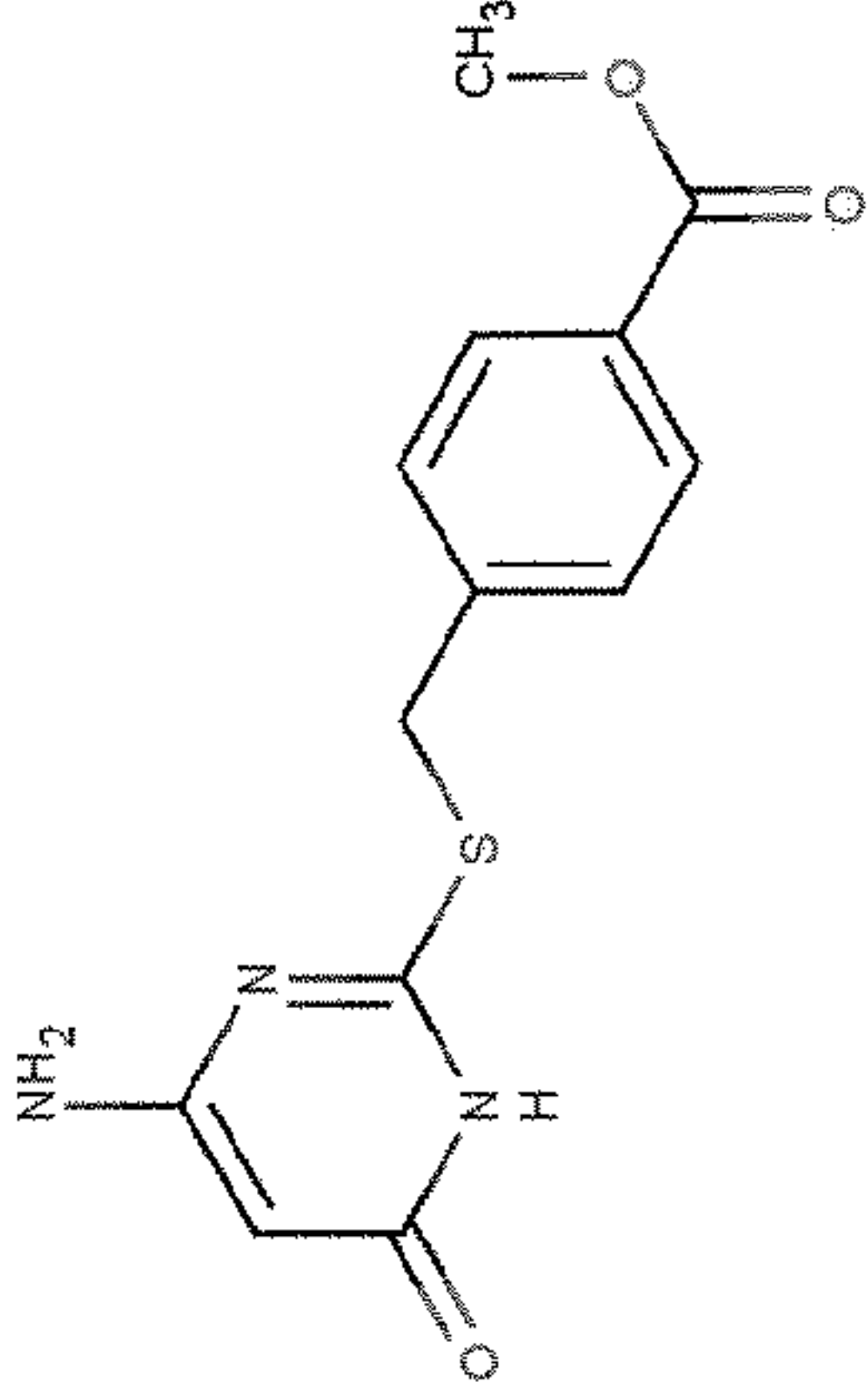
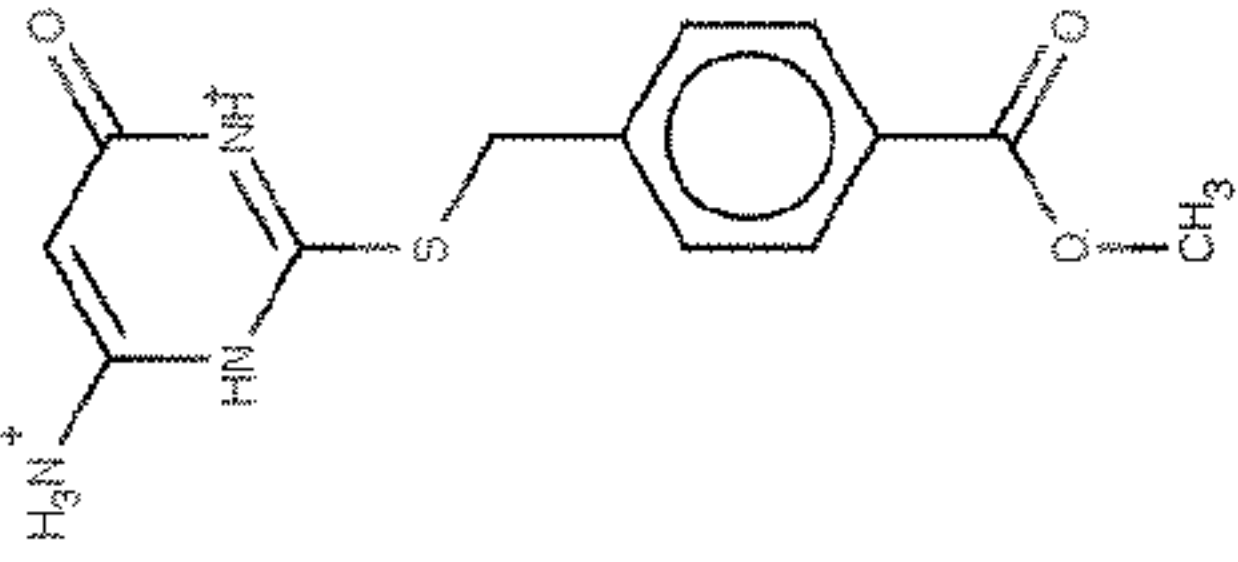
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 39		F1016-0141		F1016-0141_1_1 (1 of 2)
Compound 40		F1016-0159		F1016-0159_1_1 (1 of 2)
Compound 41		F1021-0224		F1021-0224_1_2 (1 of 4)

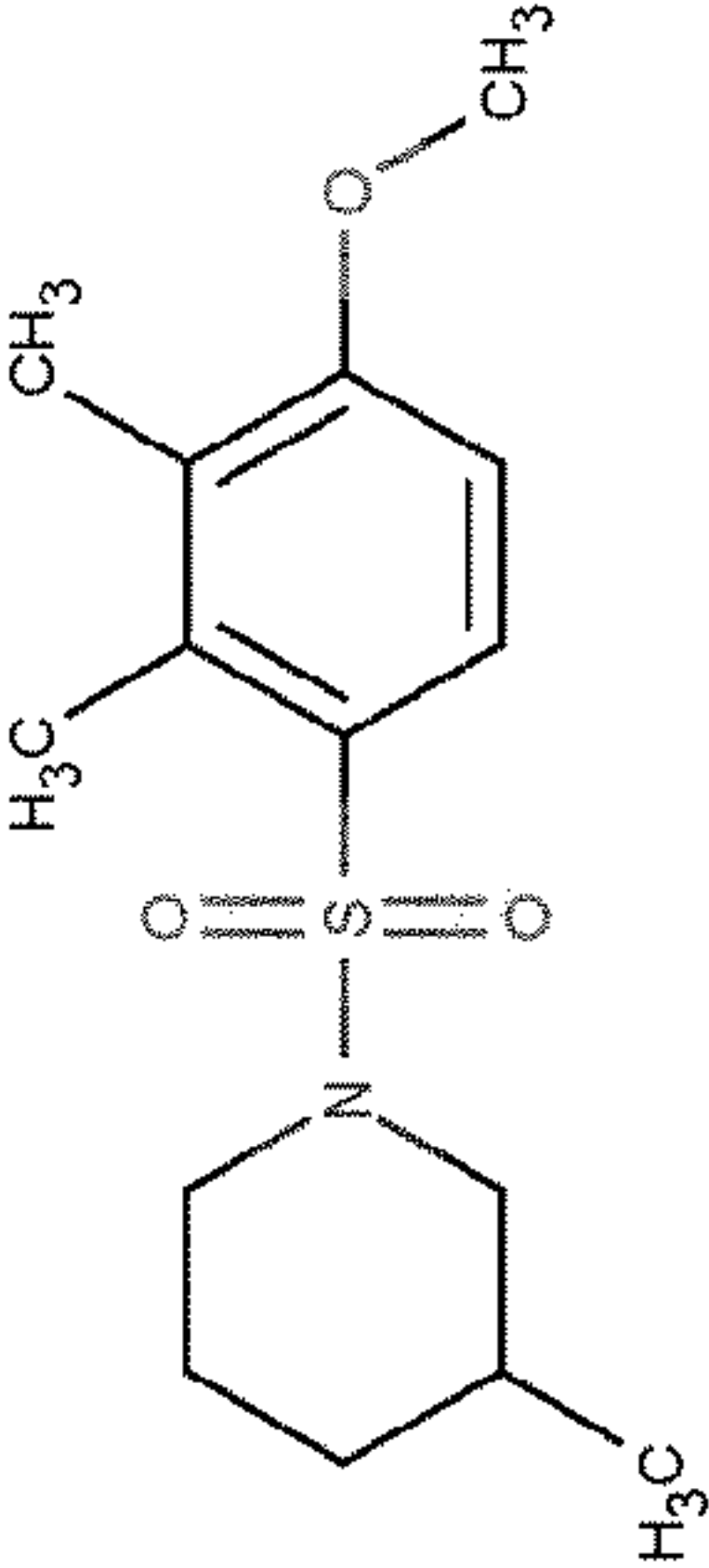
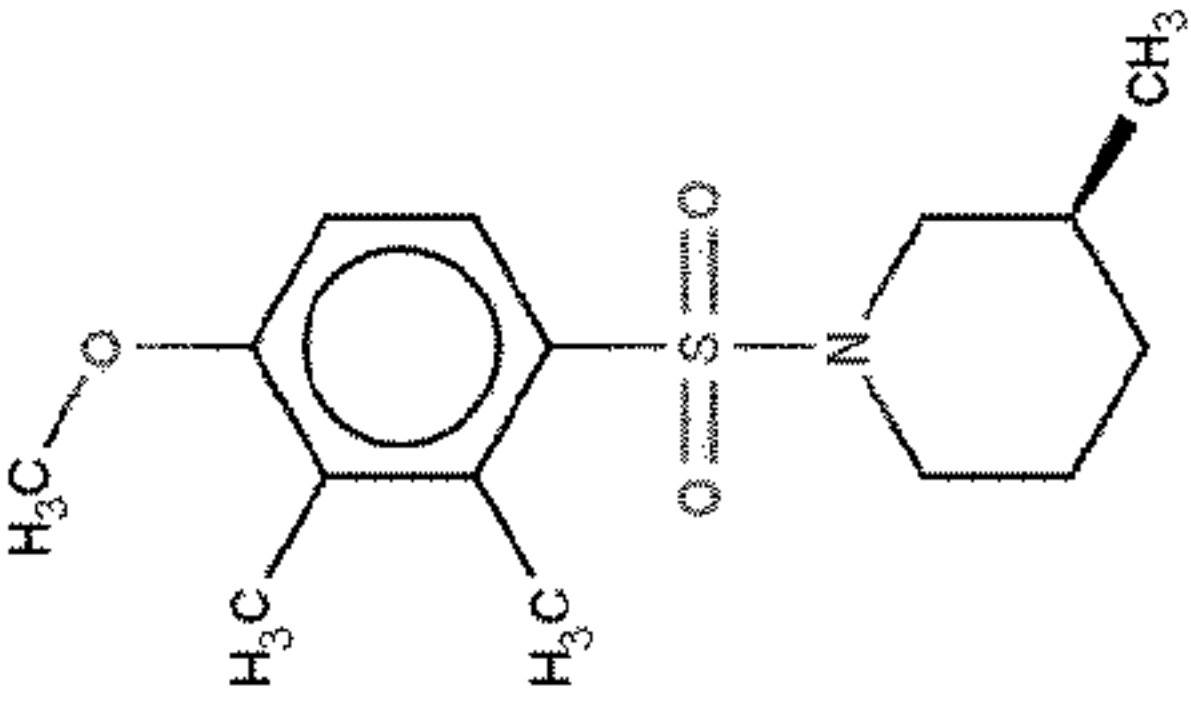
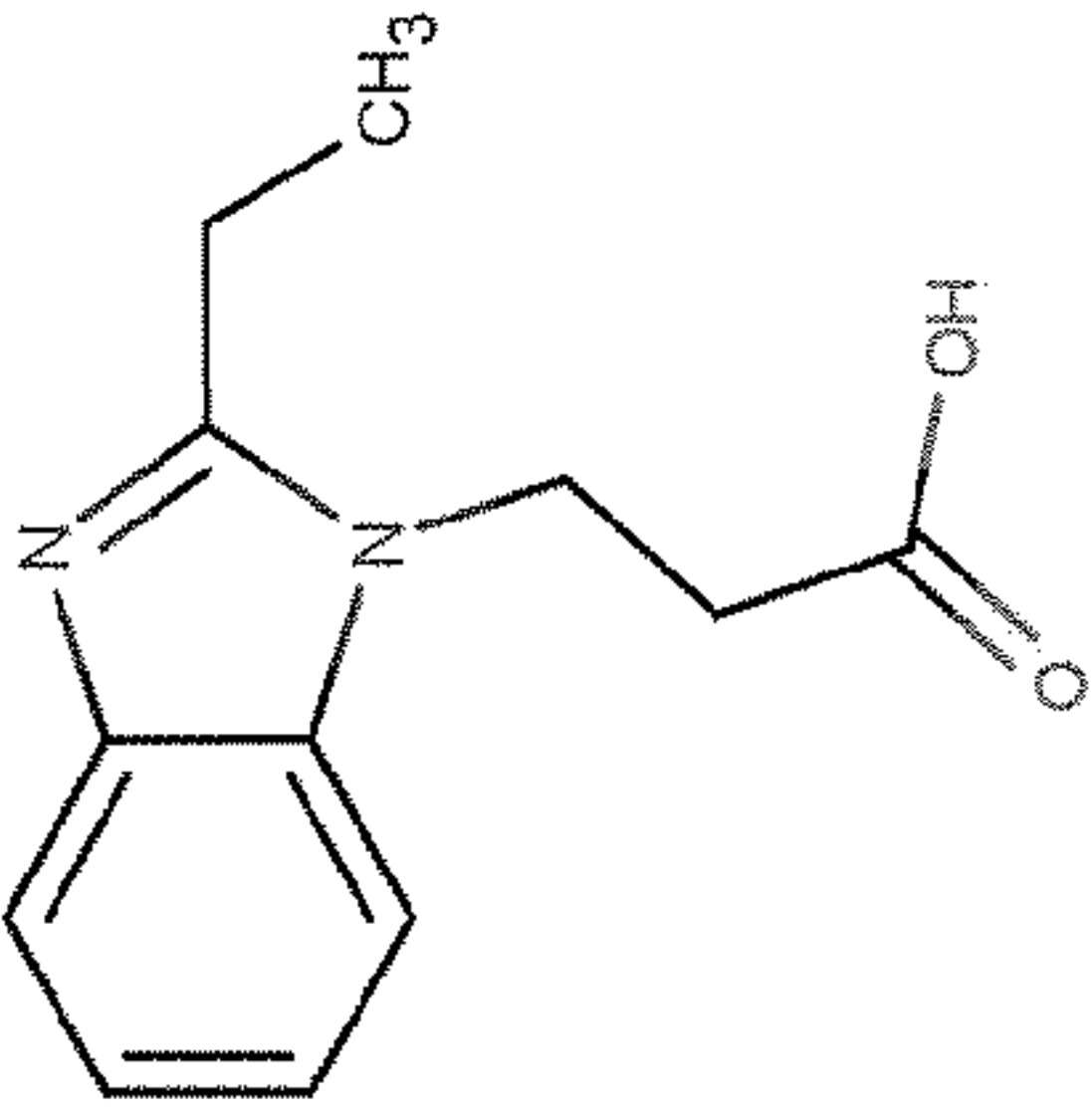
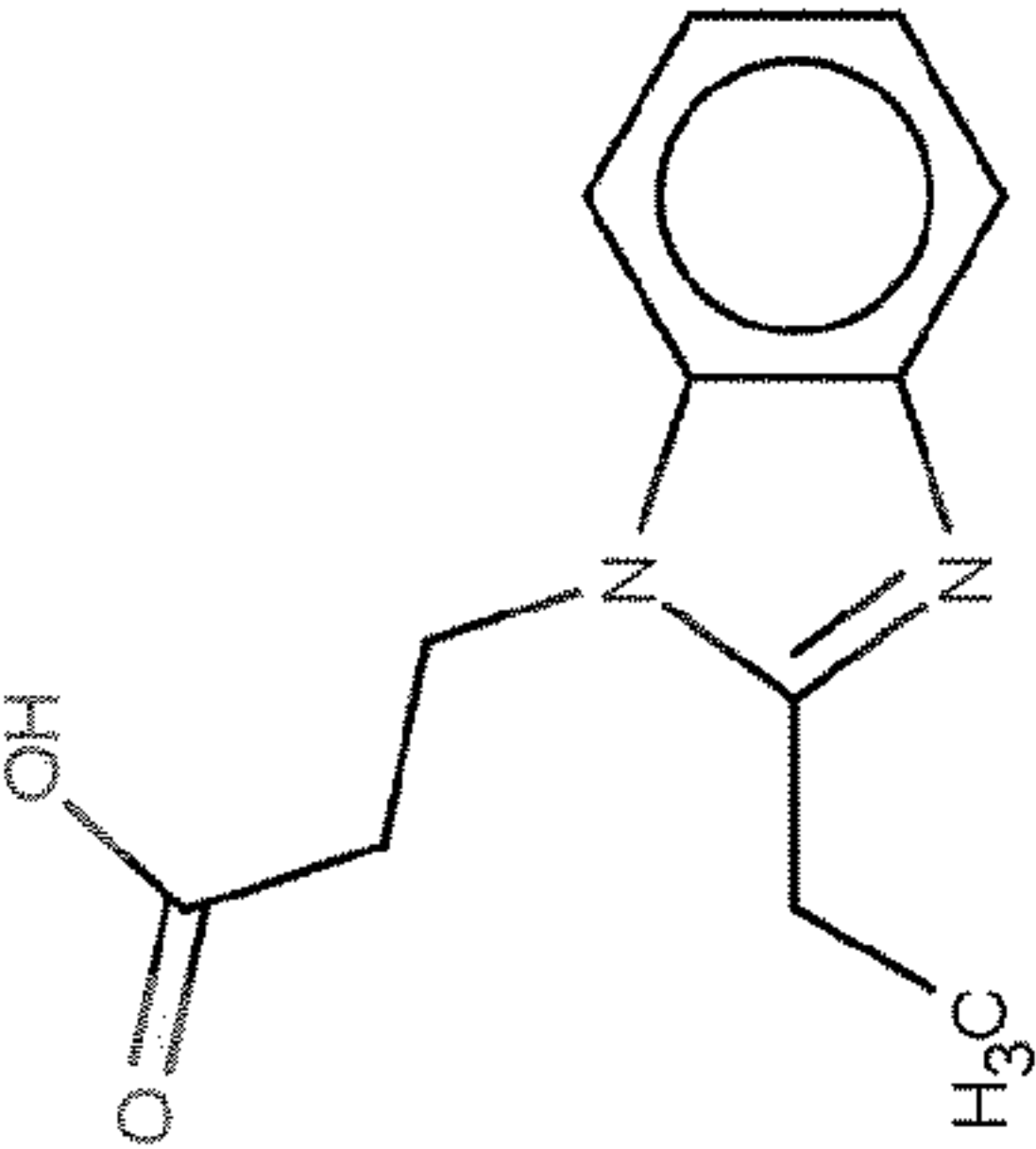
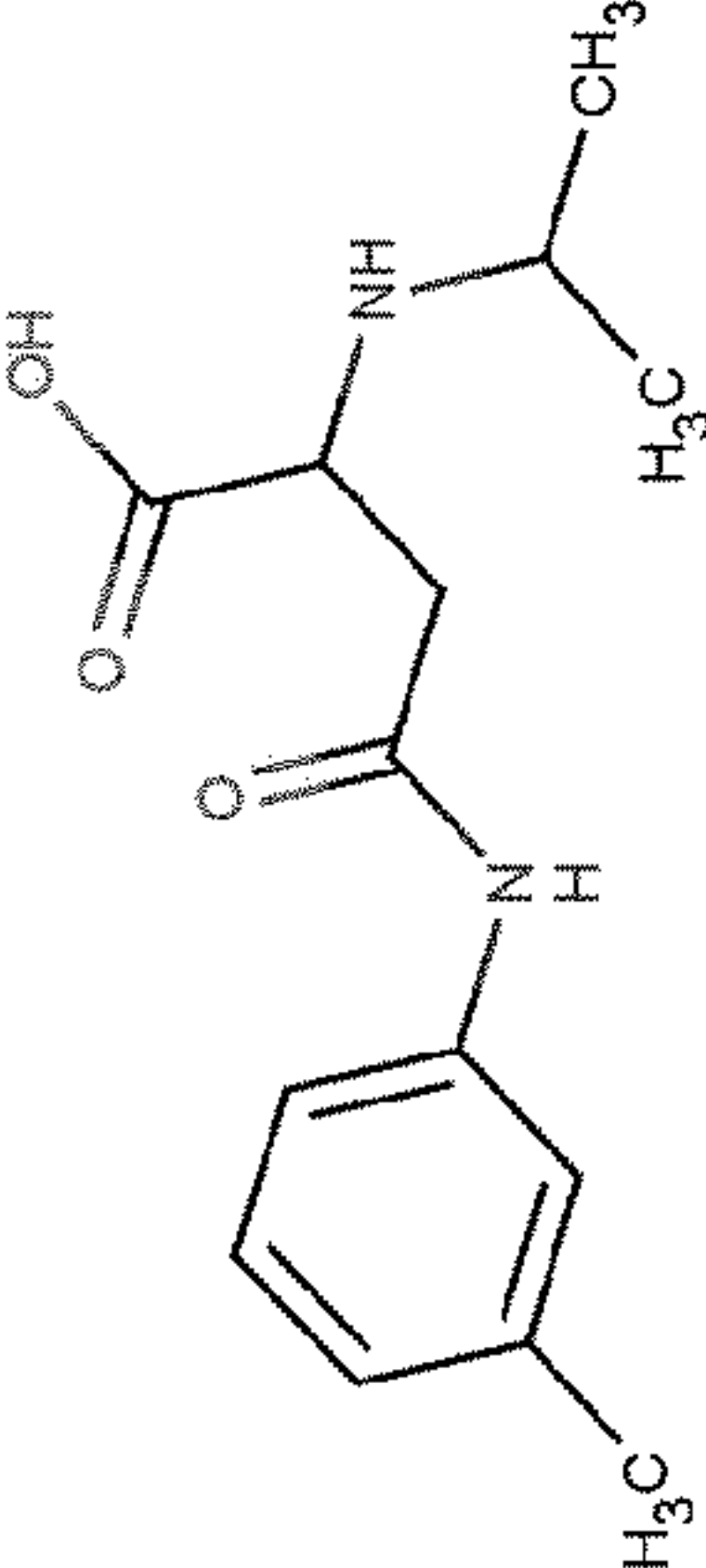
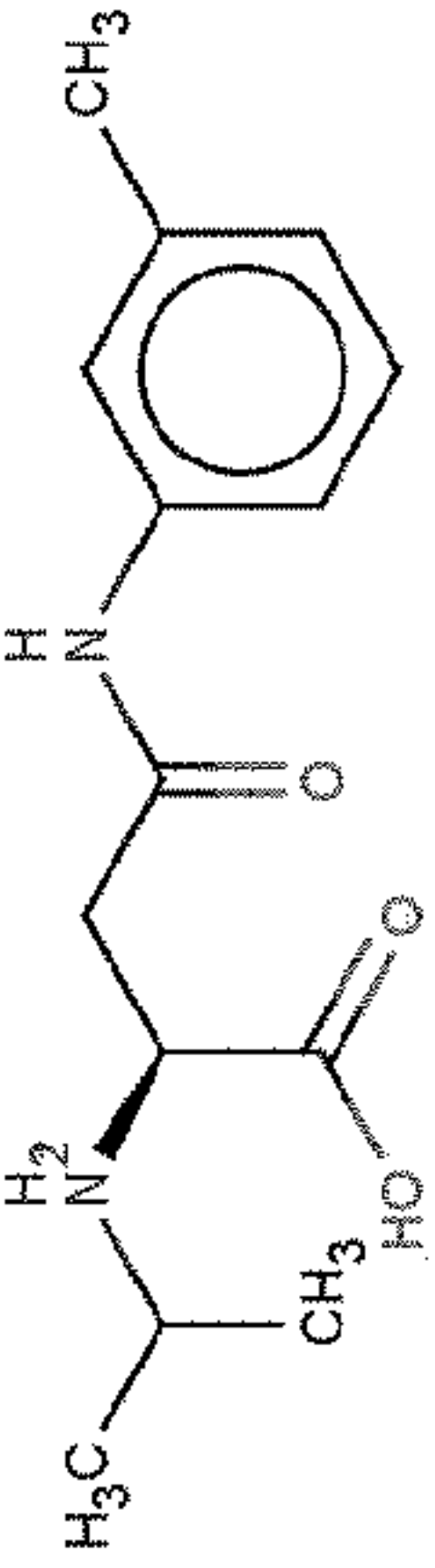
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 42		F1032-0001		F1032-0001_1_2 (1 of 4)
Compound 43		F1032-0021		F1032-0021_1_2 (1 of 4)
Compound 44		F1032-0031		F1032-0031_1 (1 of 4)

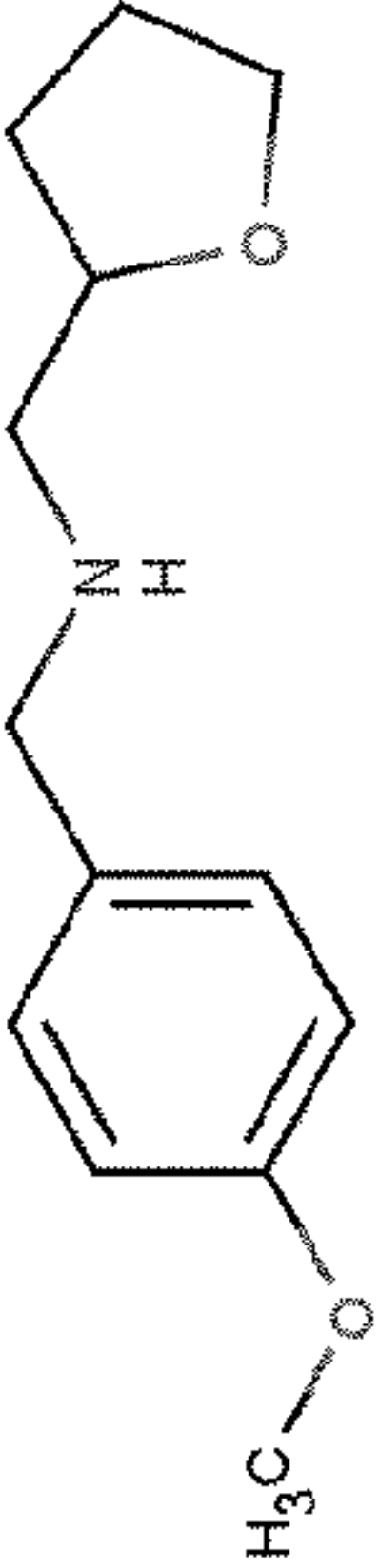
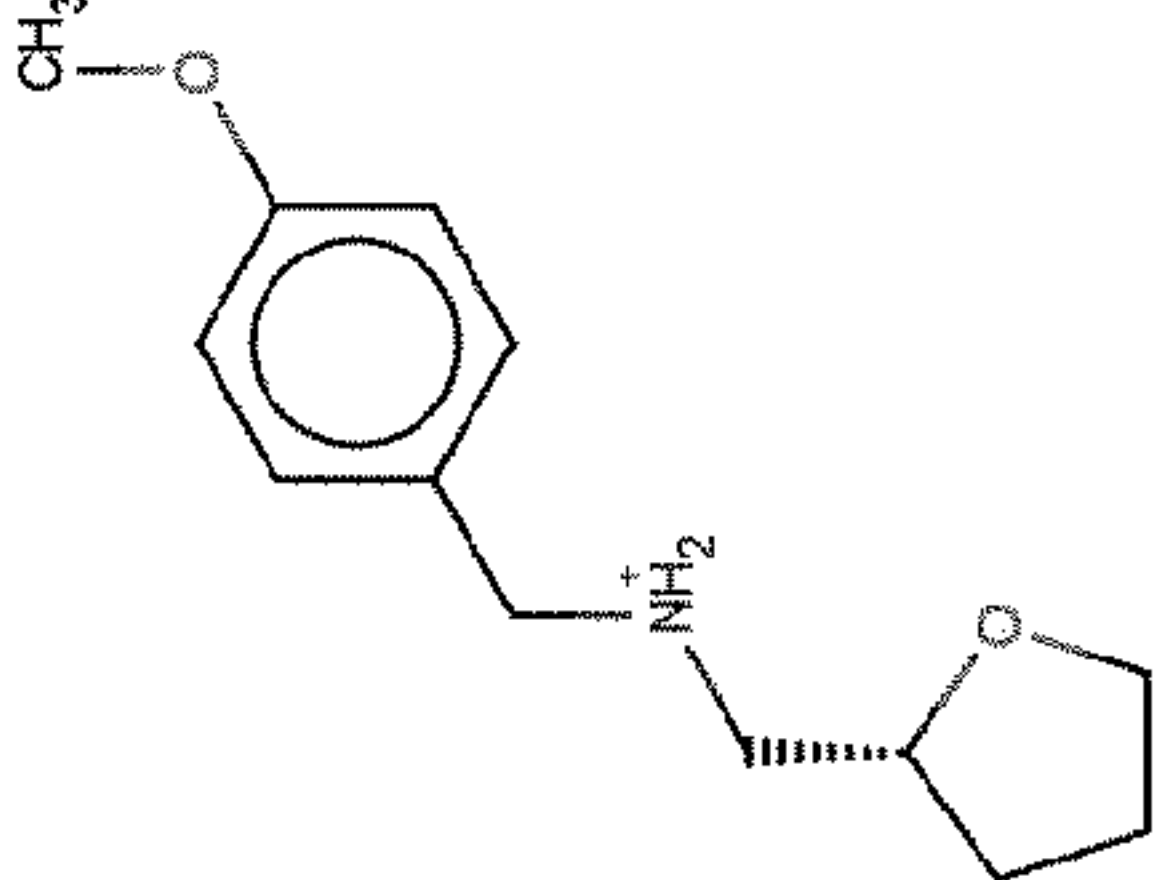
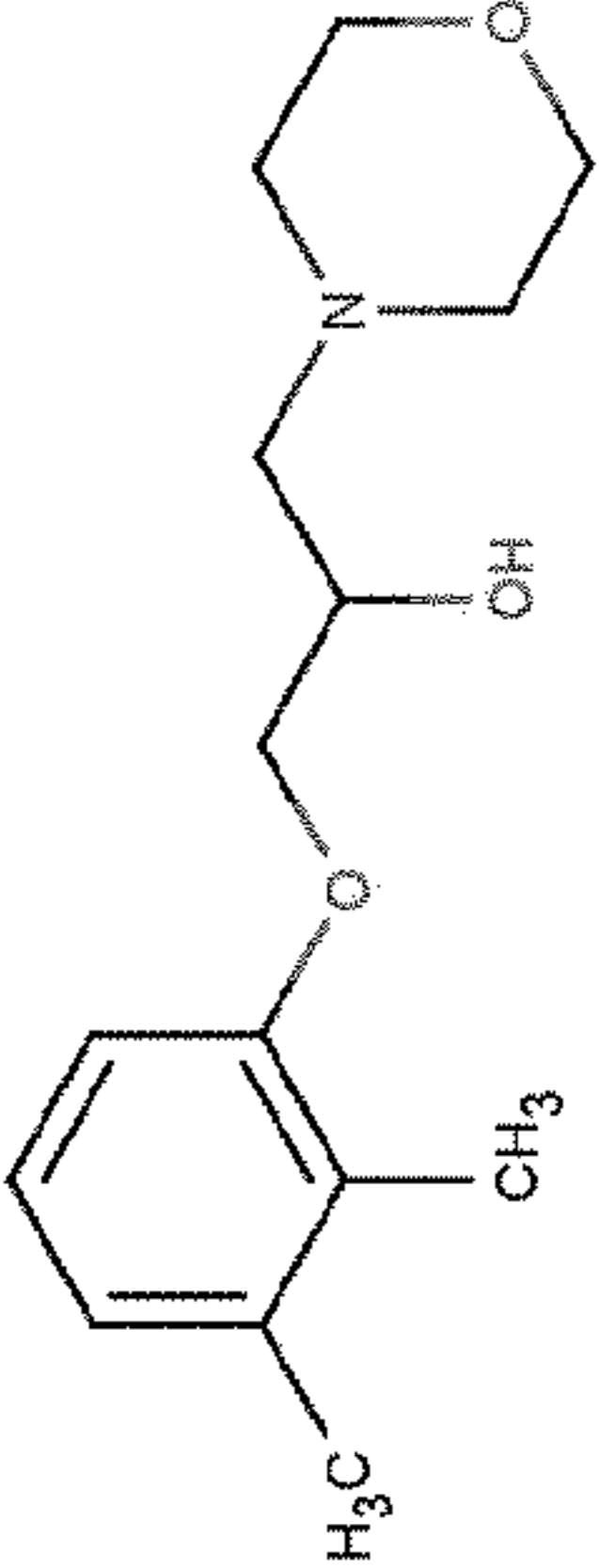
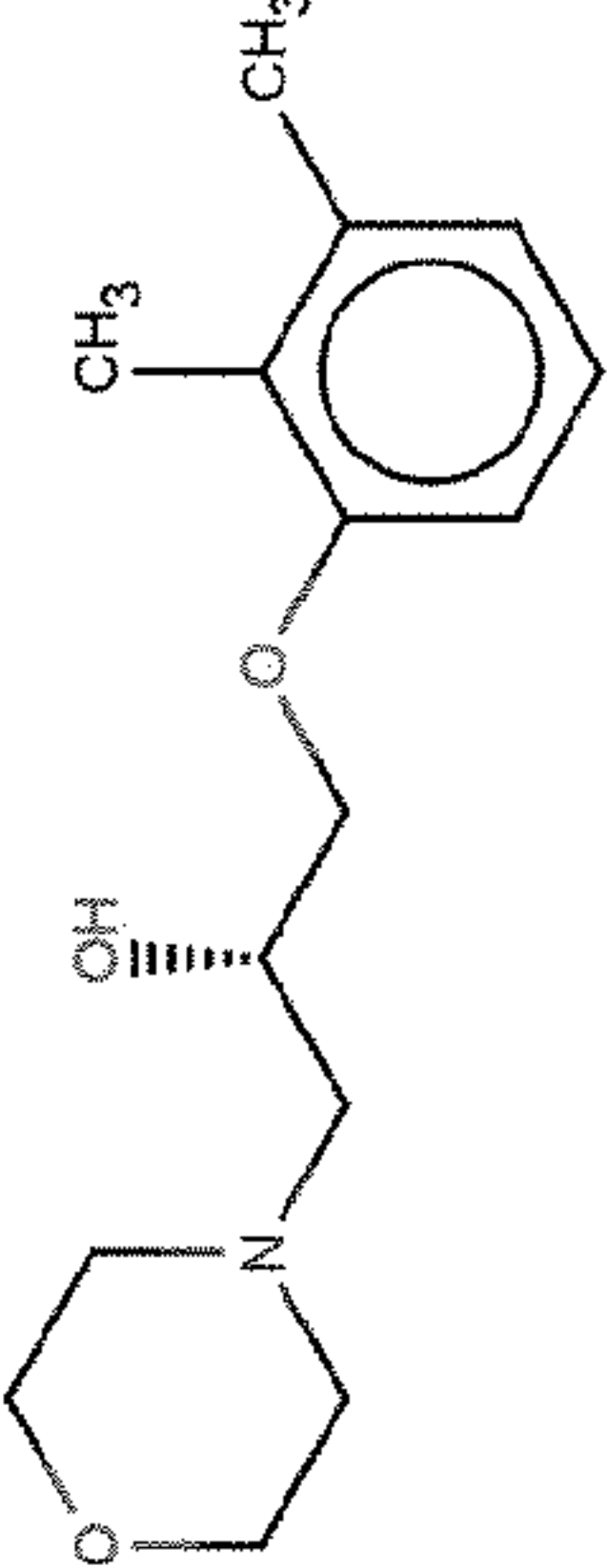
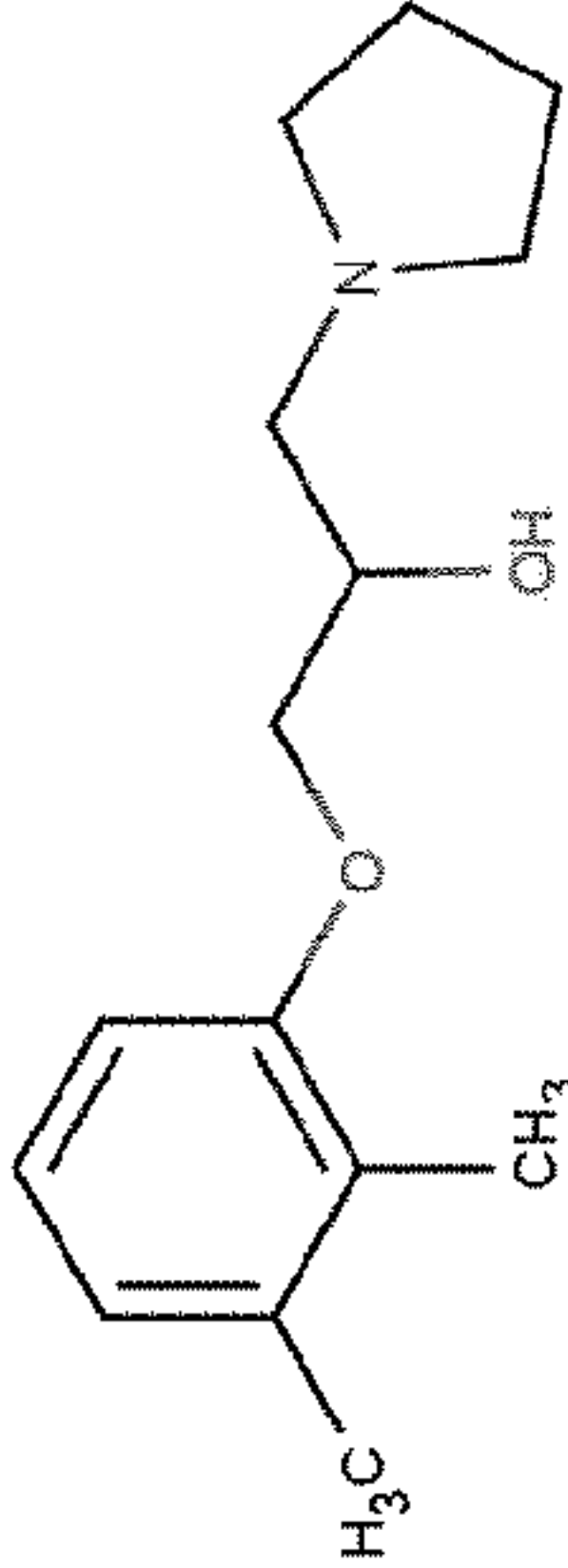
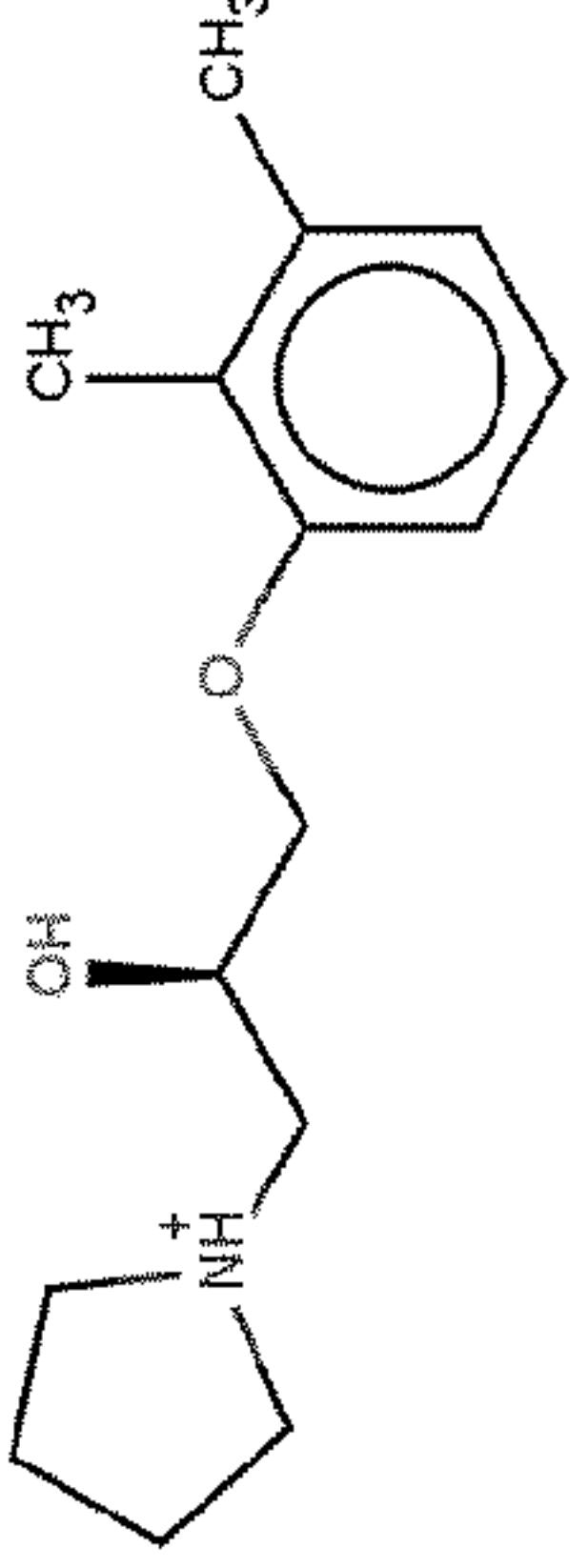
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 45		F1065-0751		F1065-0751_3_1 (1 of 8)
Compound 46		F1107-0031		F1107-0031_80_4 (1 of 448)
Compound 47		F1126-0569		F1126-0569_4_1 (1 of 6)

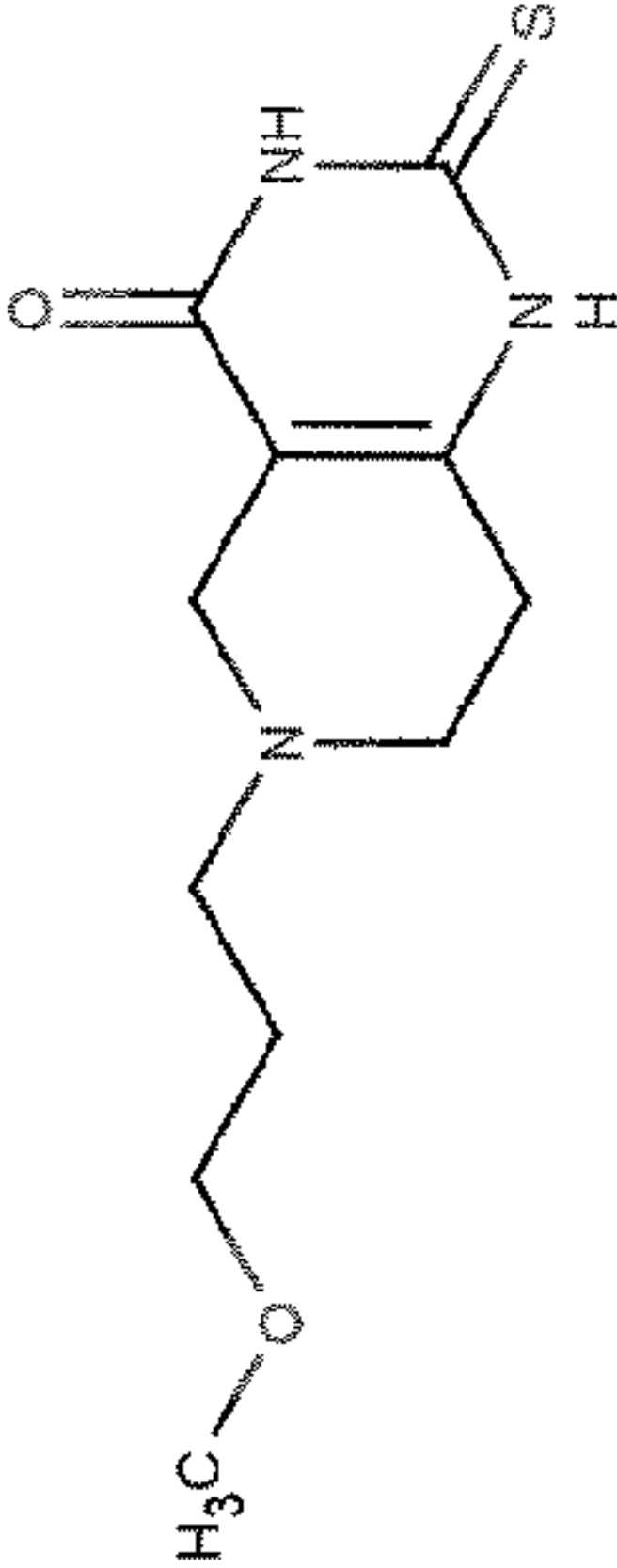
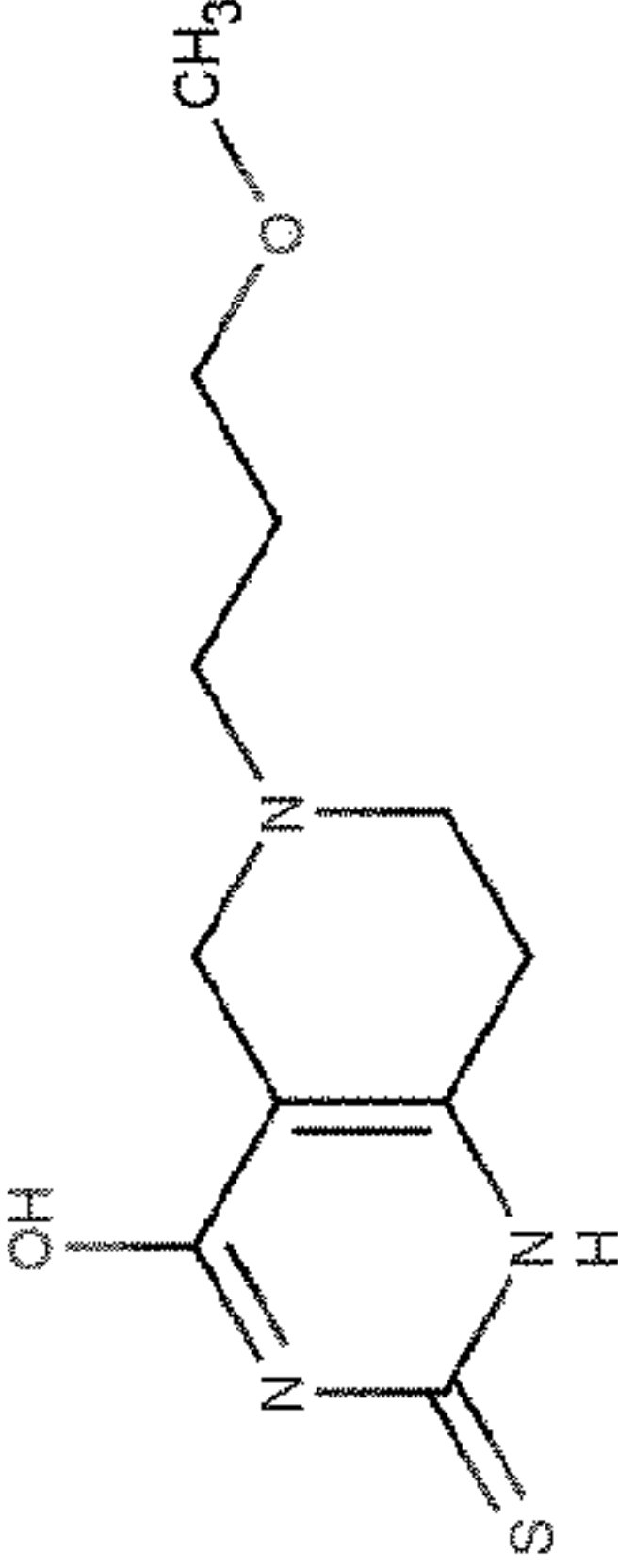
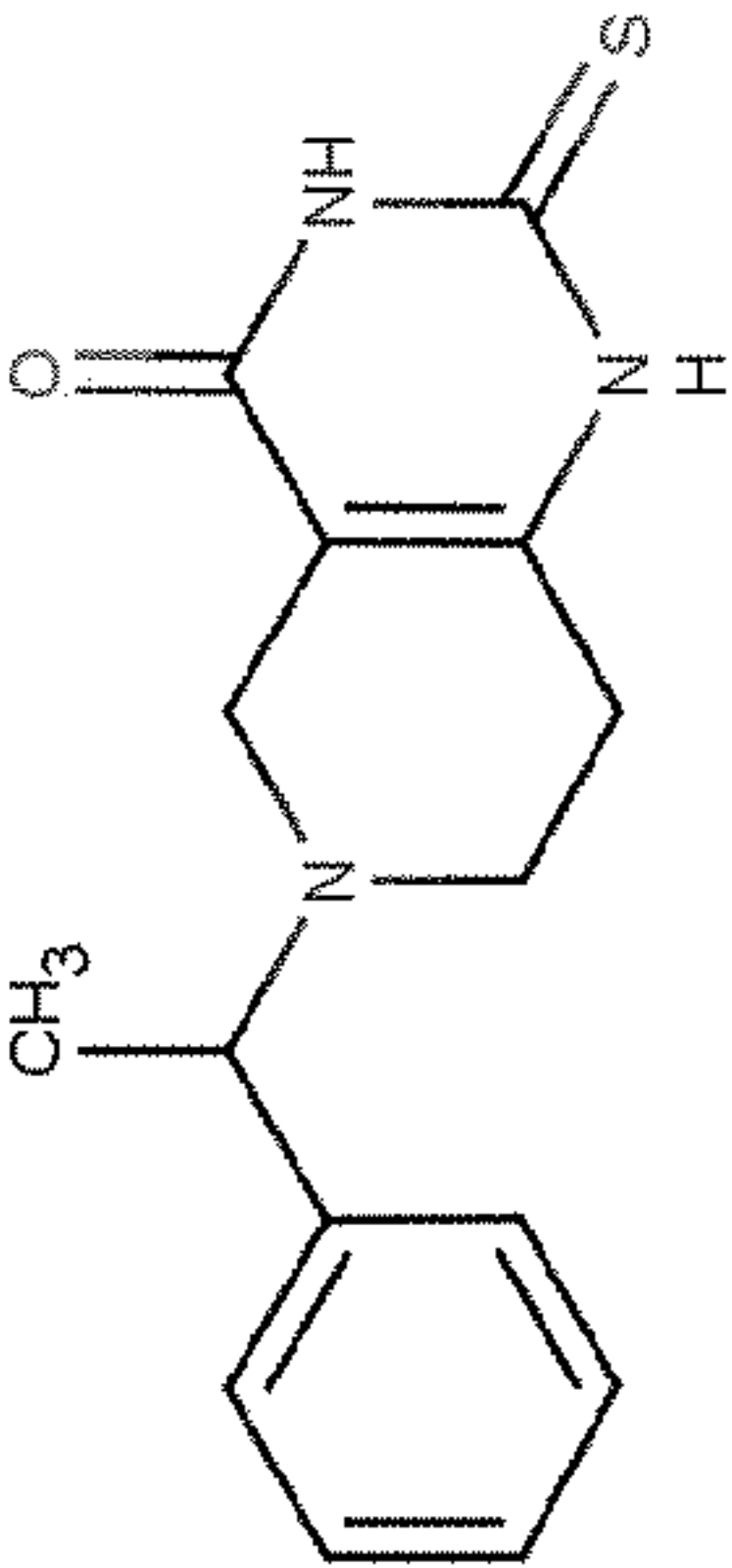
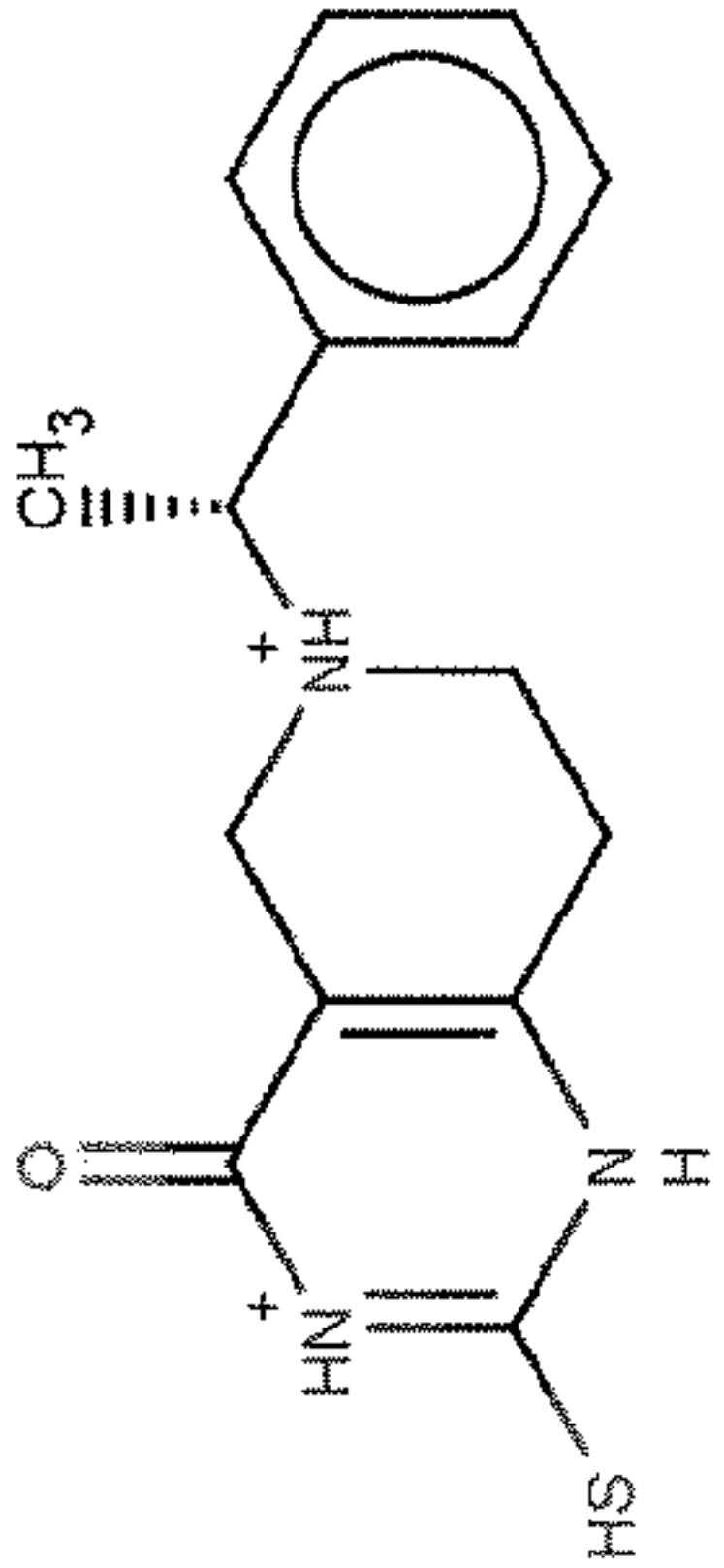
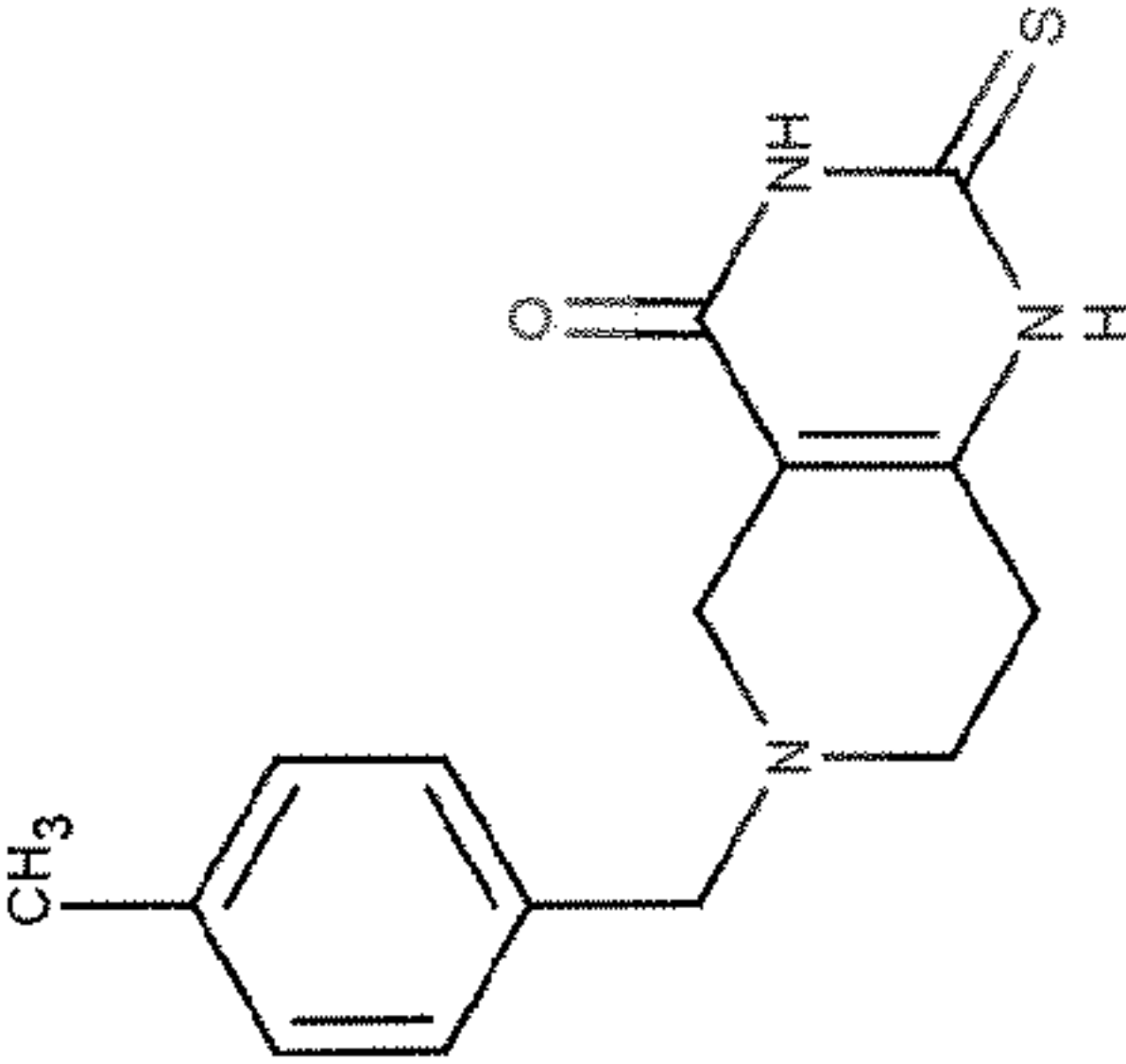
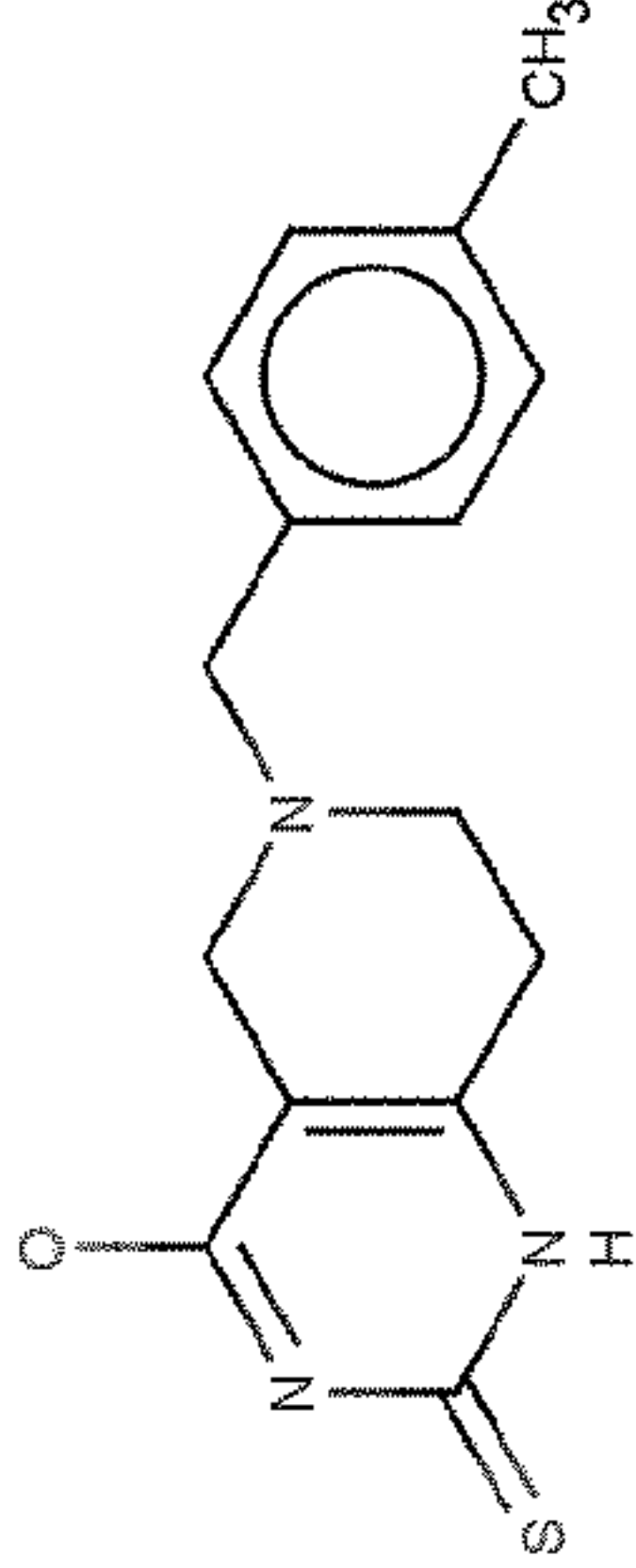
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 48		F1132-0388		F1132-0388_1_1 (1 of 1)
Compound 49		F1174-1011		F1174-1011_1_1 (1 of 2)
Compound 50		F1243-0186		F1243-0186_3_1 (1 of 6)

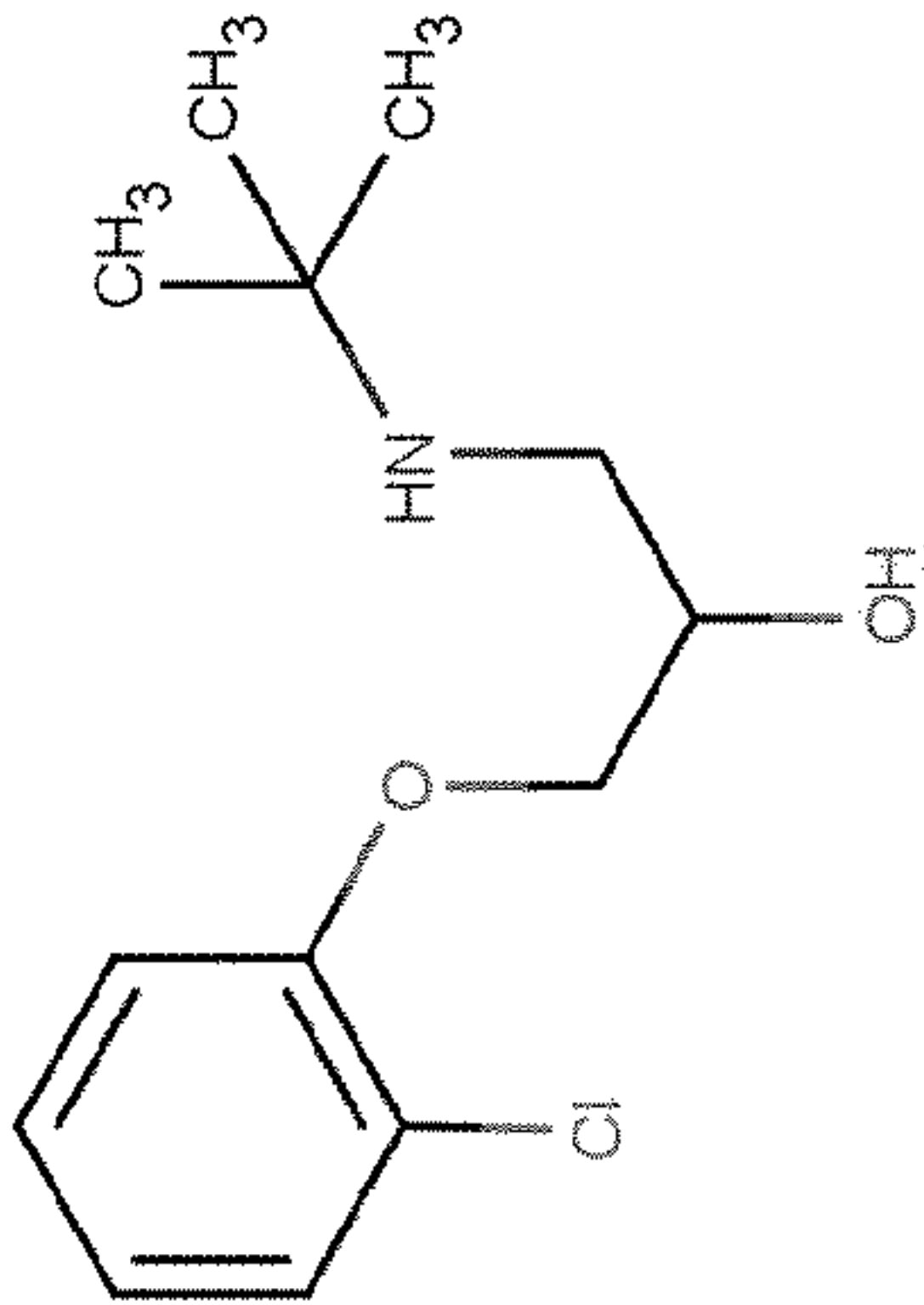
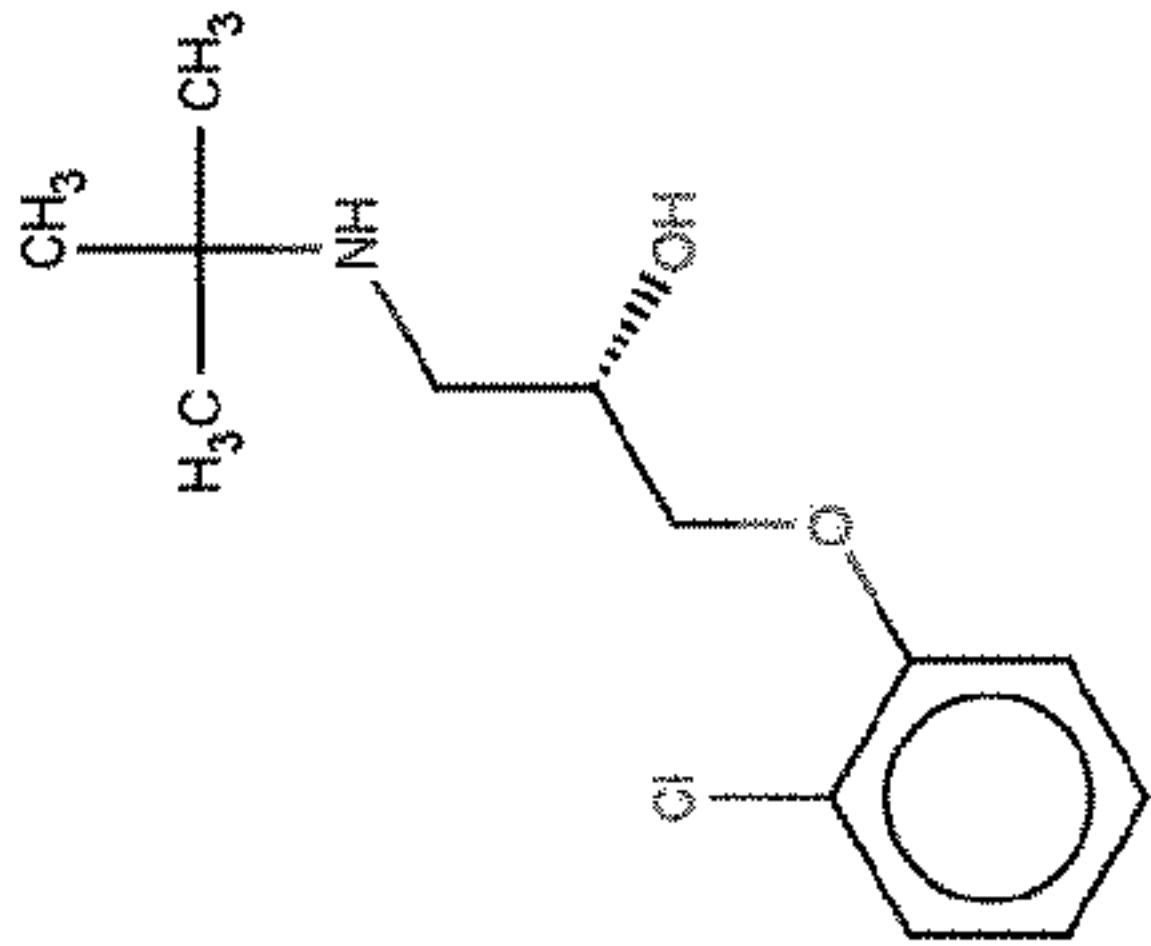
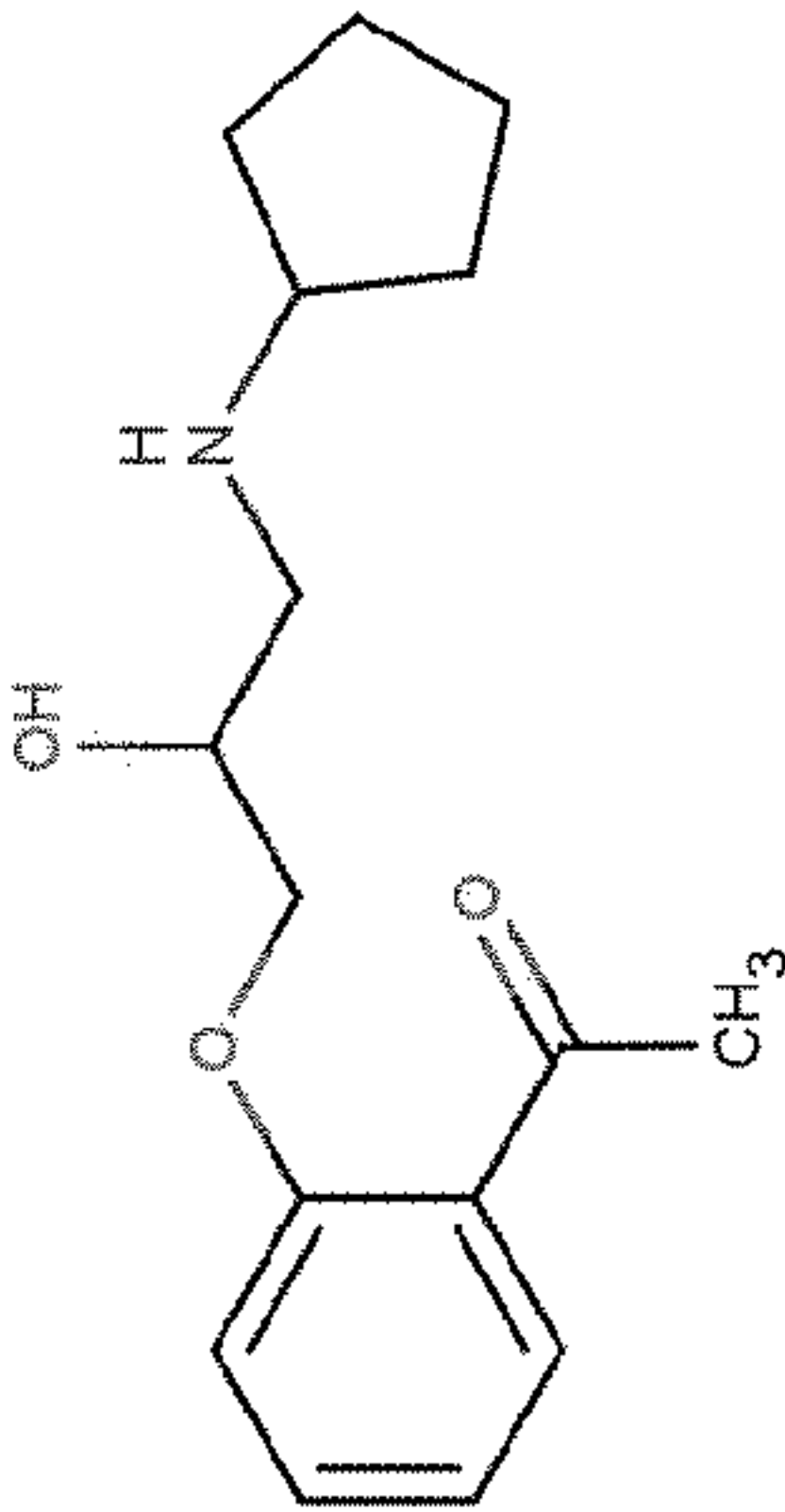
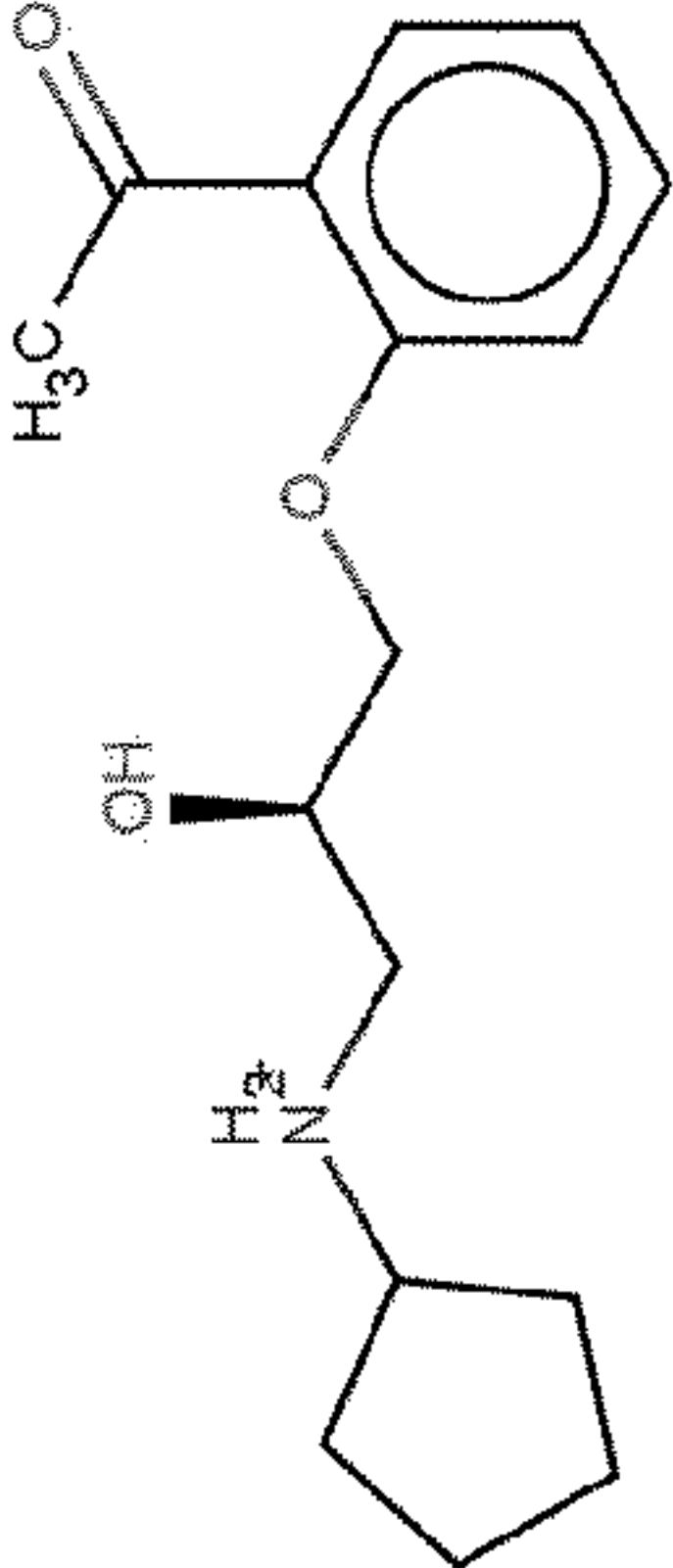
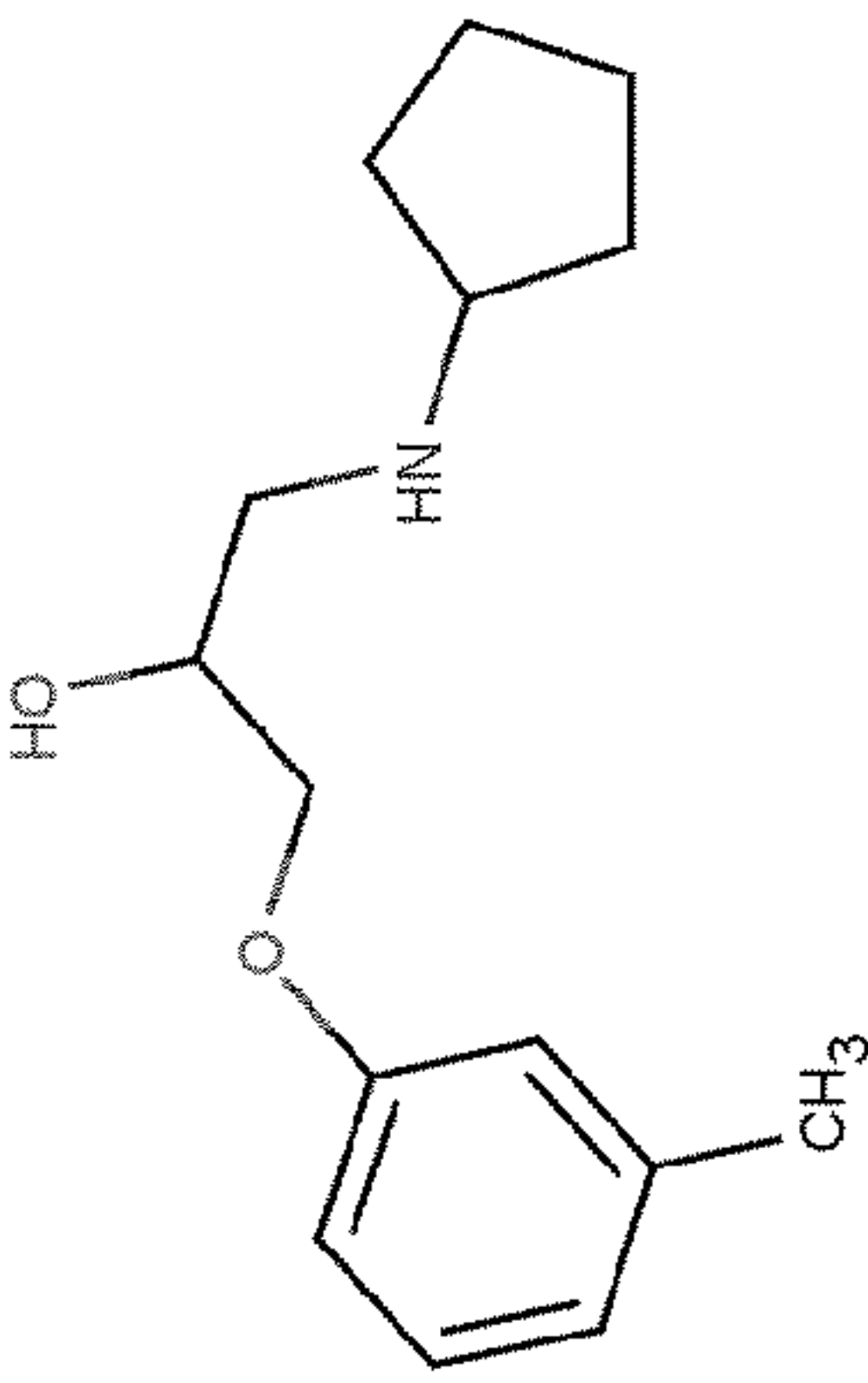
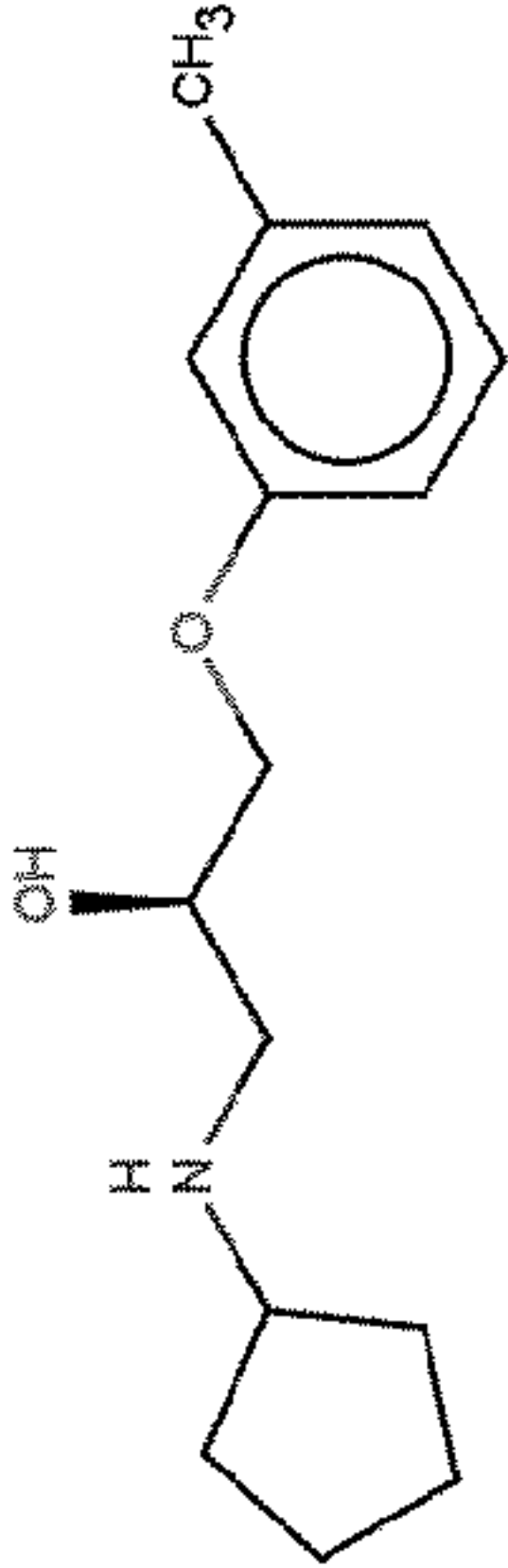
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 51		F1360-0381		F1360-0381_3_1 (1 of 4)
Compound 52		F1387-0270		F1387-0270_2_1 (1 of 3)
Compound 53		F1411-0031		F1411-0031_9_1 (1 of 14)

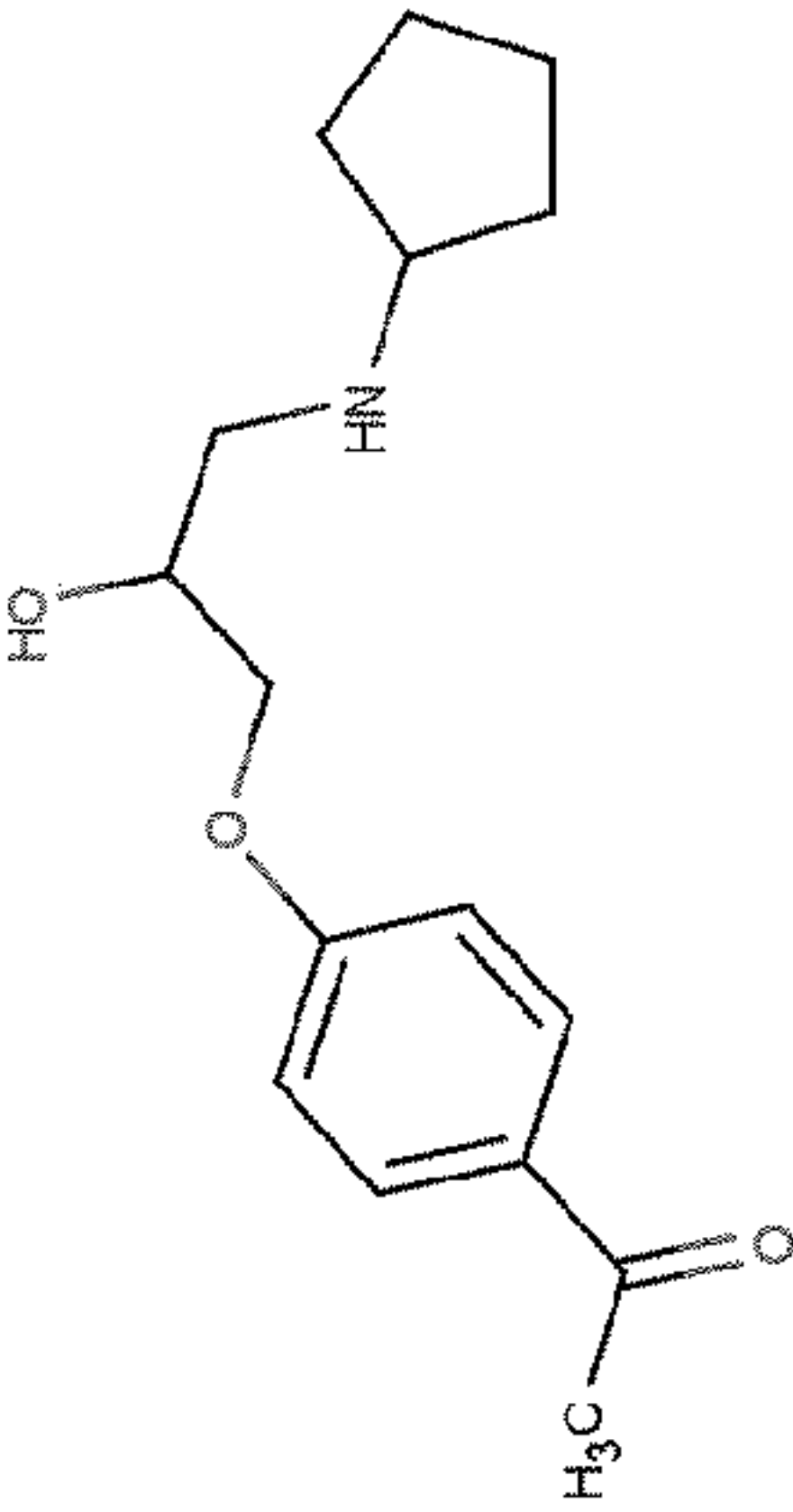
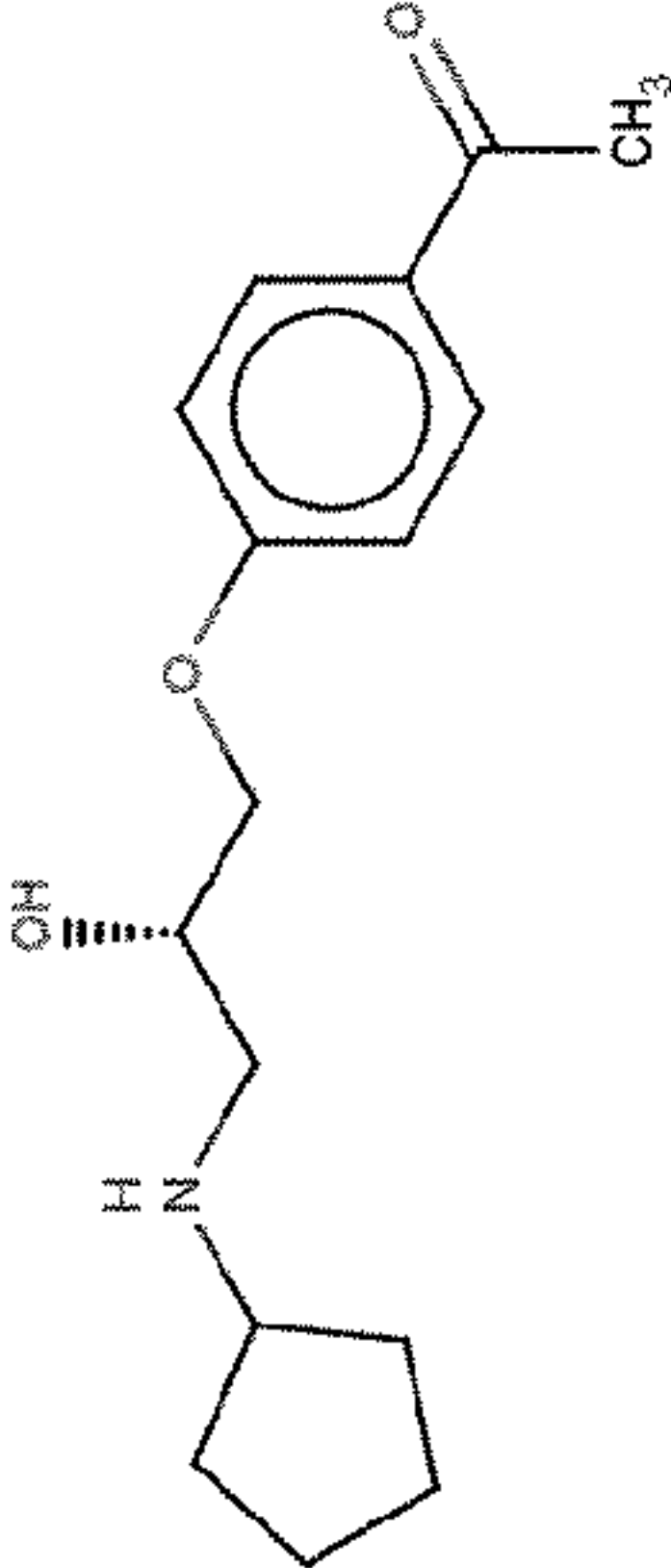
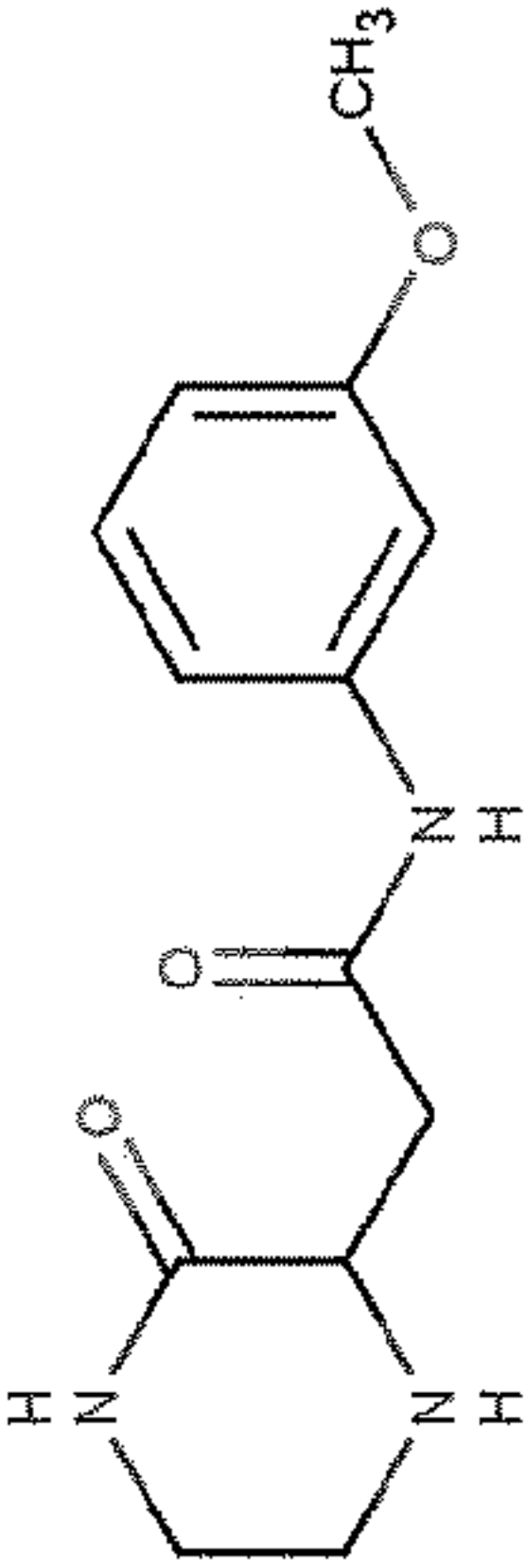
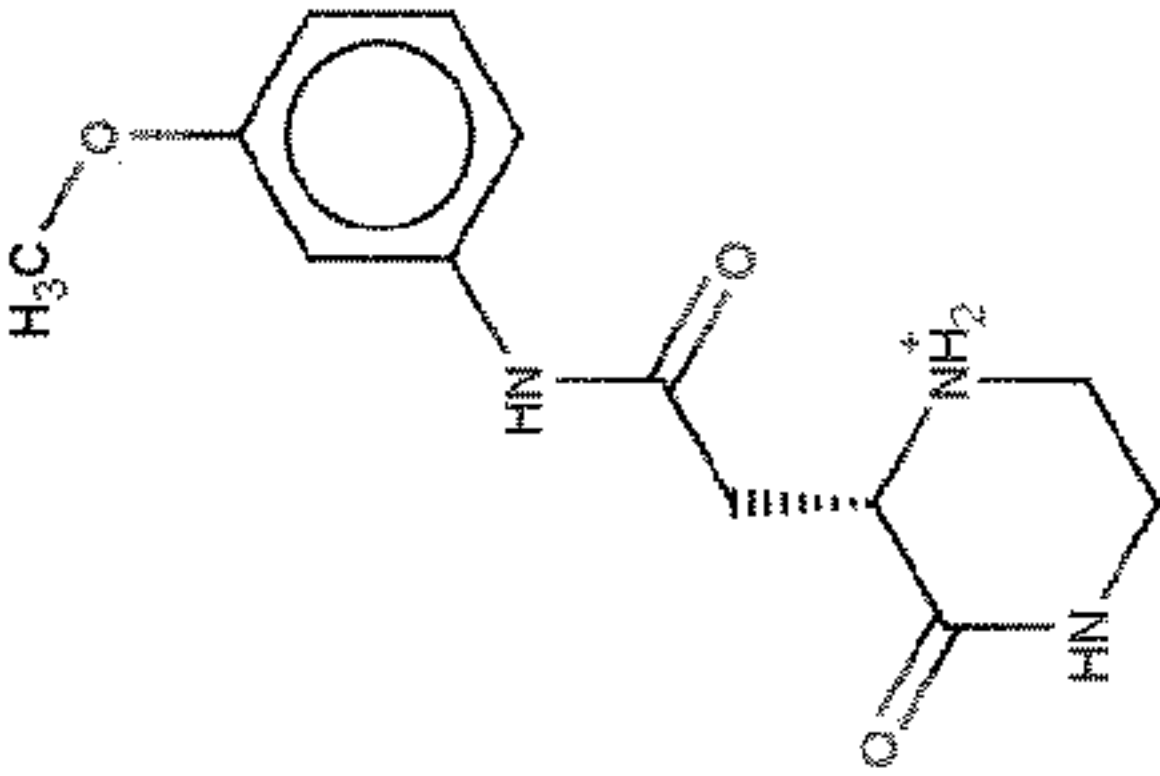
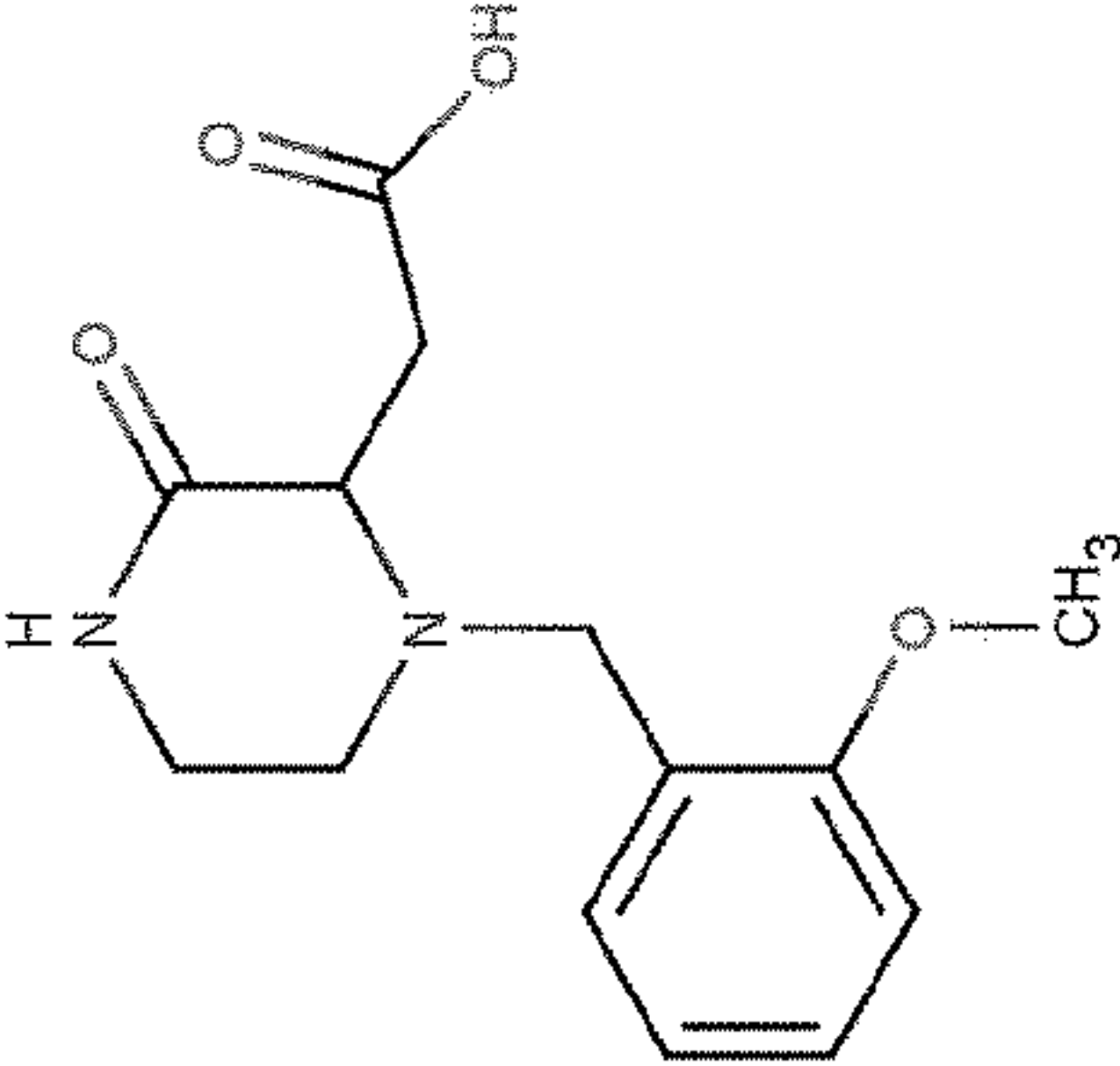
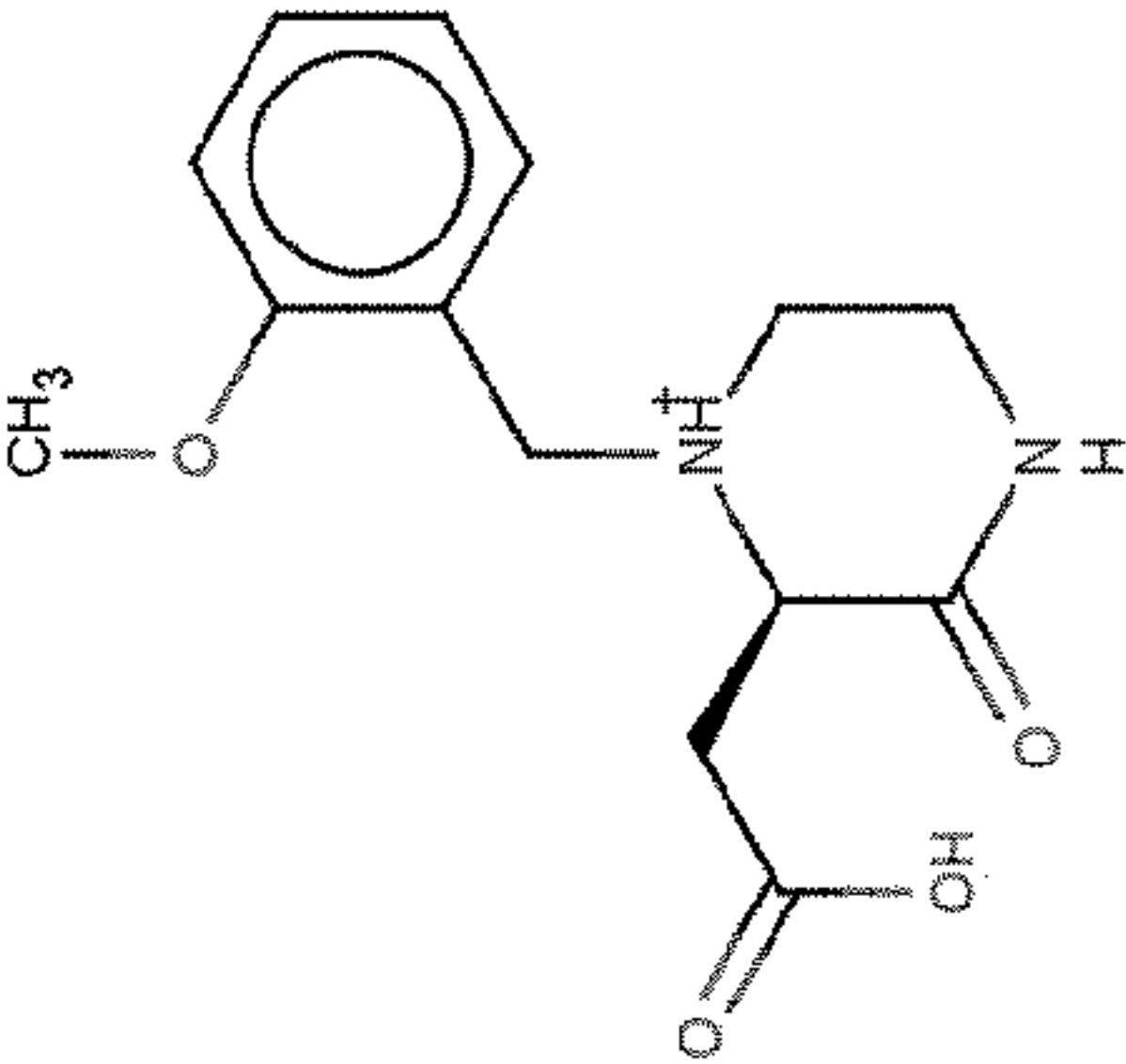
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 54		F1438-0002		F1438-0002_2 (1 of 4)
Compound 55		F1438-0006		F1438-0006_1_3 (1 of 8)
Compound 56		F1440-0010		F1440-0010_9_1 (1 of 14)

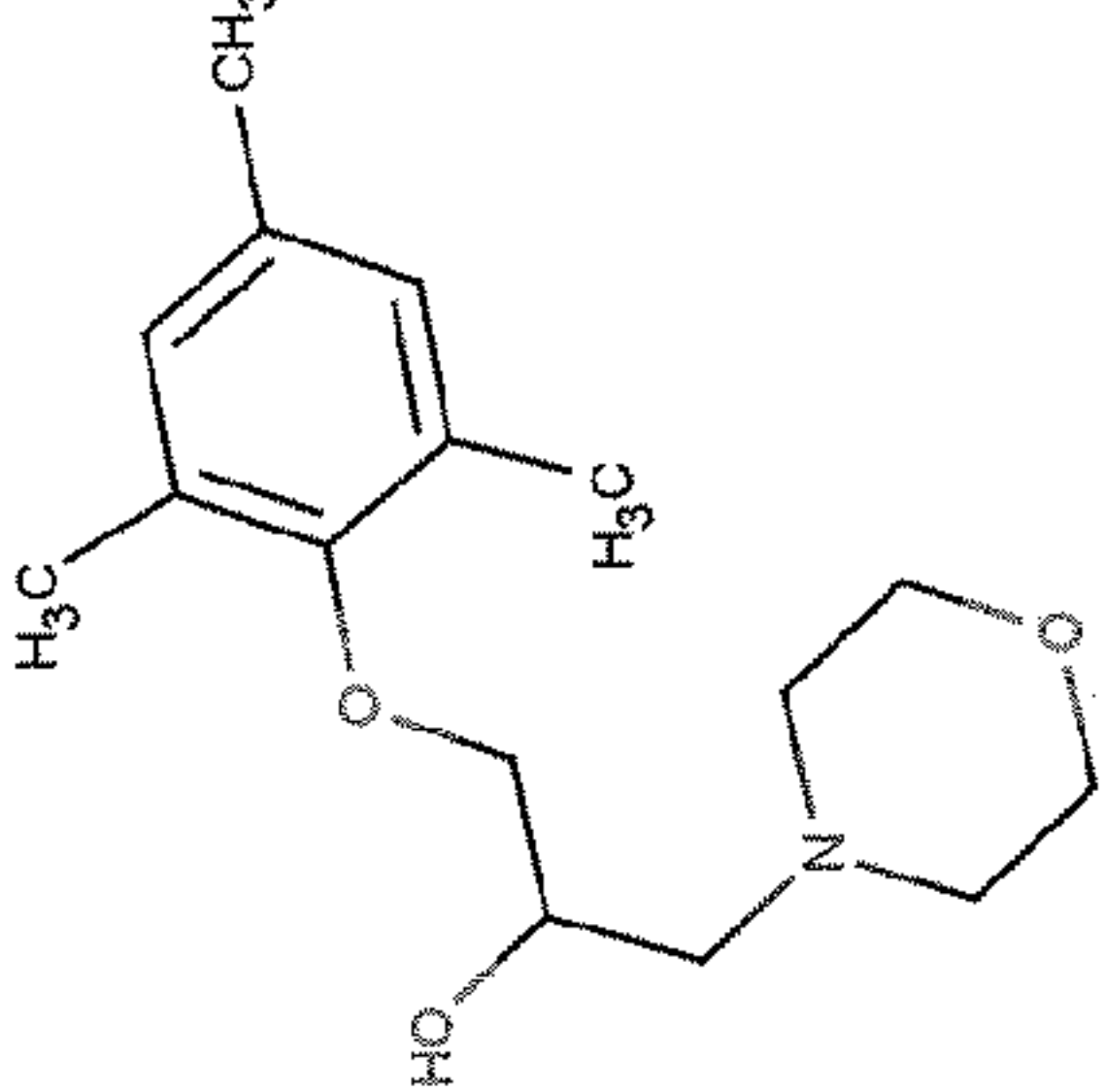
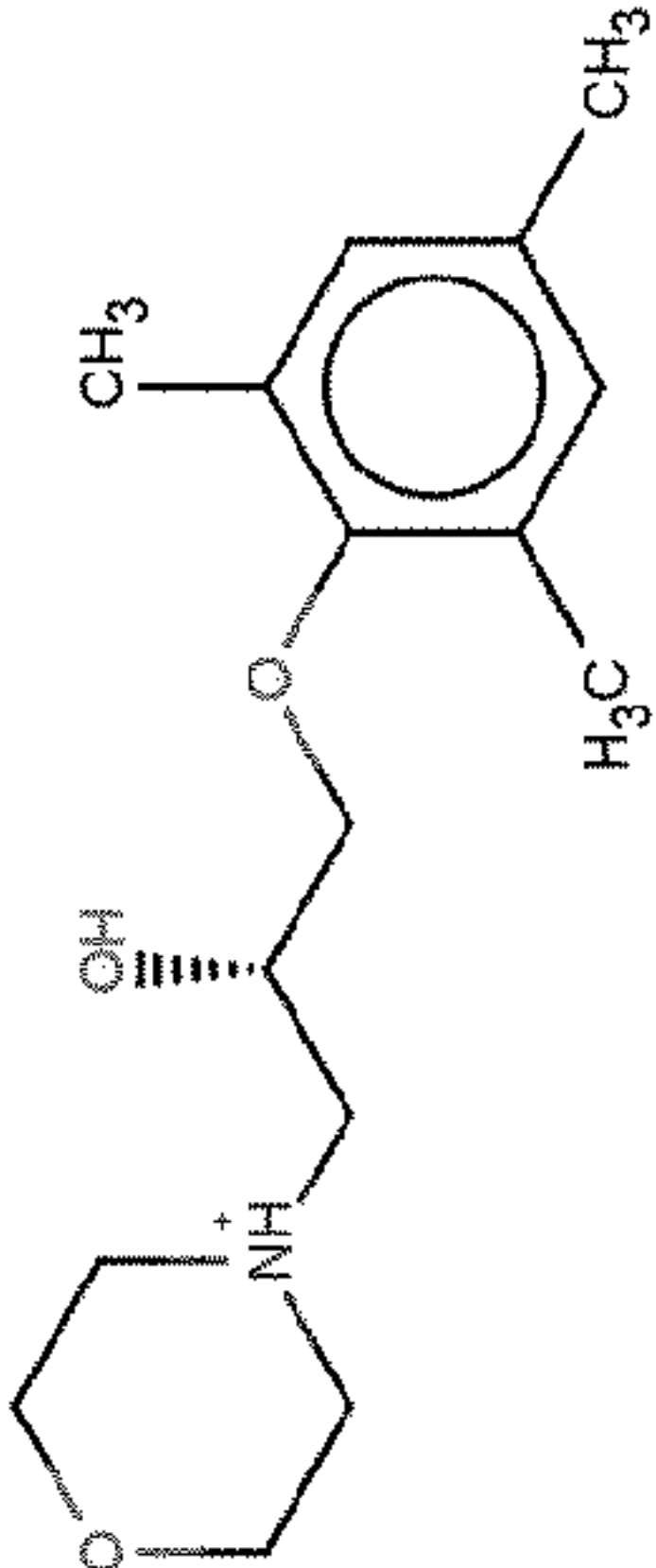
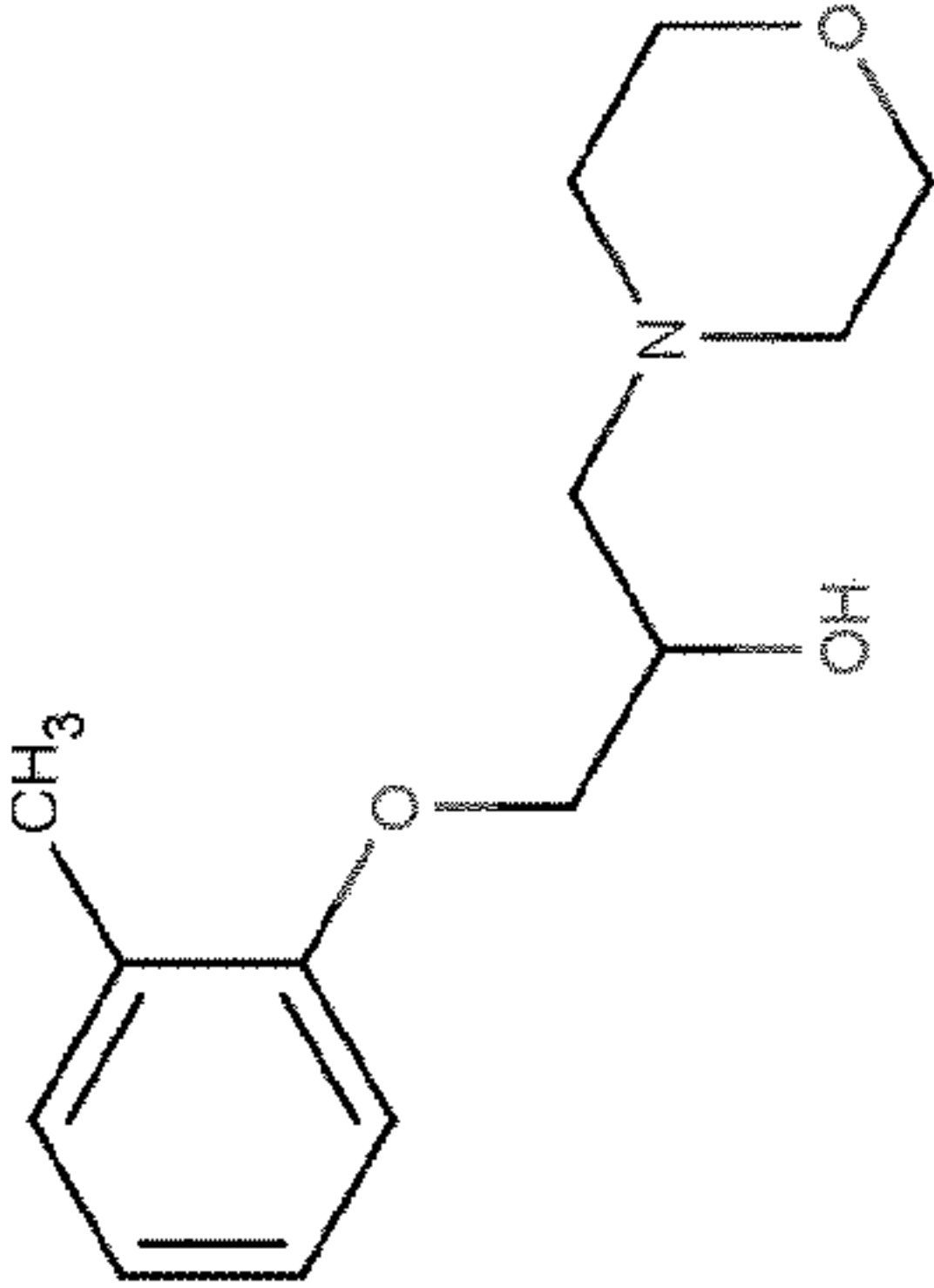
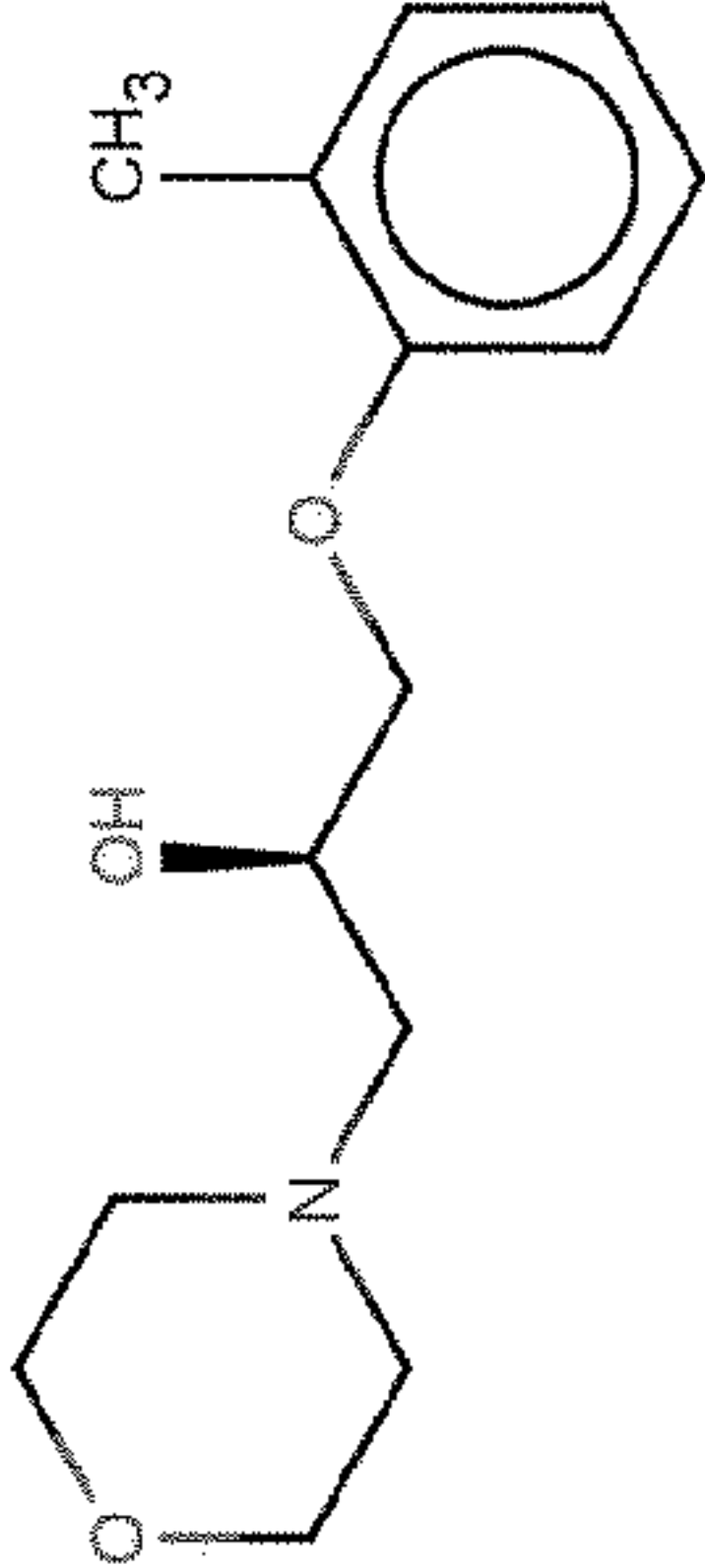
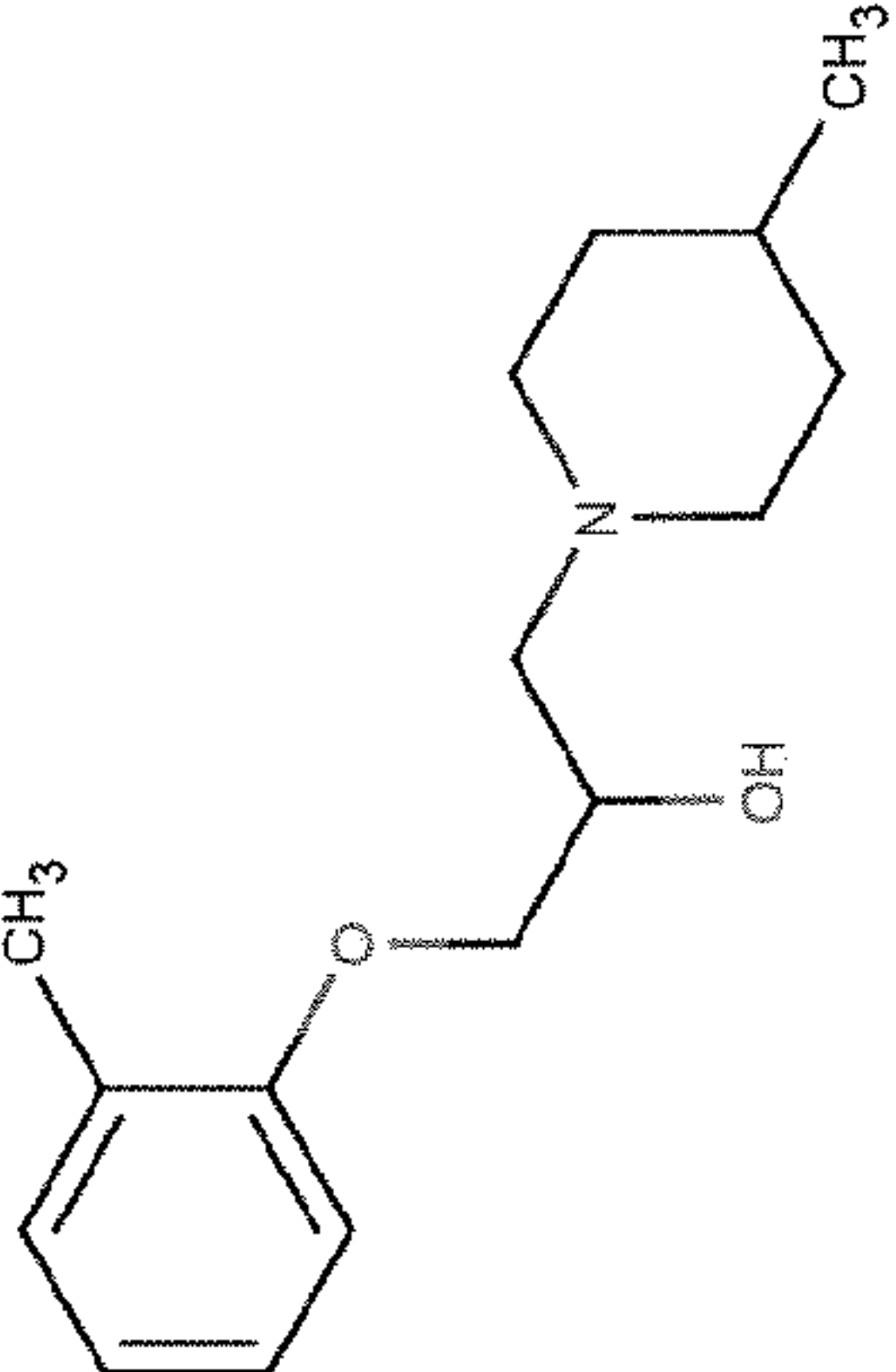
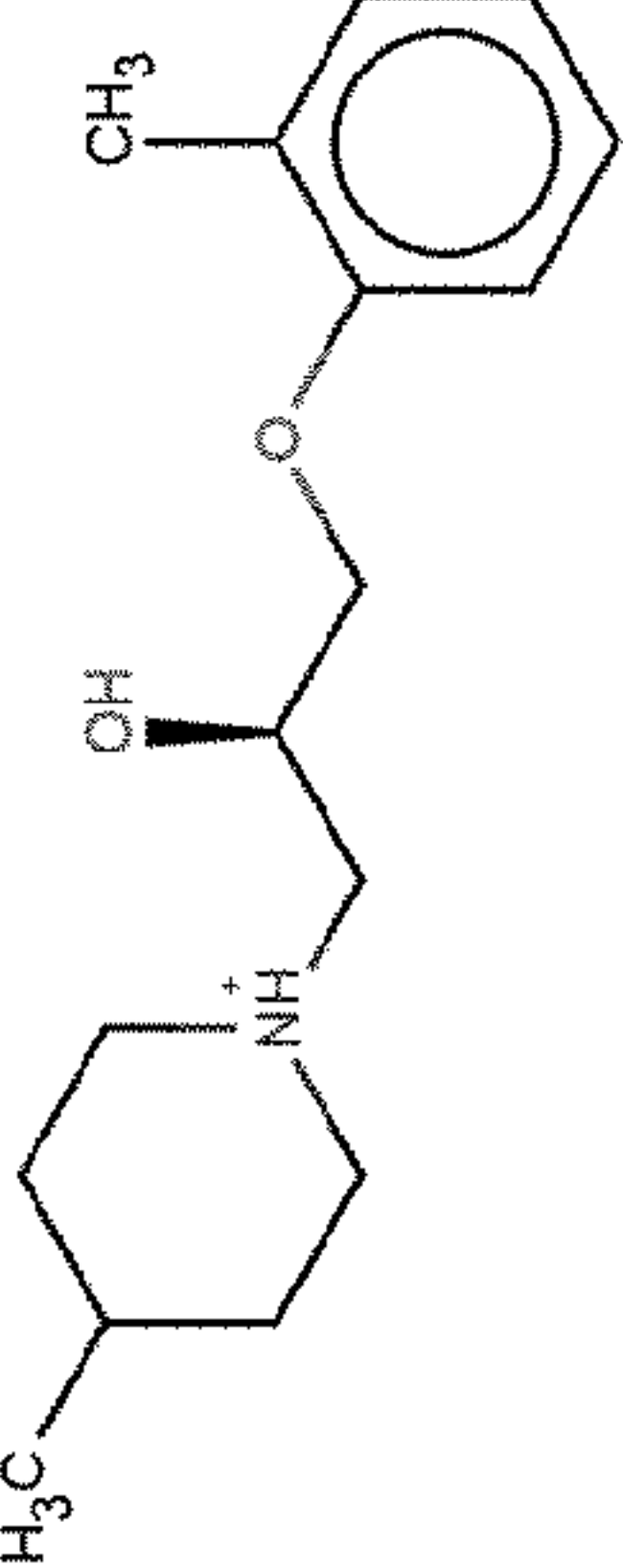
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 57		F1478-0152		F1478-0152_1 (1 of 2)
Compound 58		F1501-0026		F1501-0026_1 (1 of 4)
Compound 59		F1533-0056		F1533-0056_2_2 (1 of 8)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 60		F1549-0007		F1549-0007_1_1 (1 of 4)
Compound 61		F1562-0050		F1562-0050_1 (1 of 4)
Compound 62		F1562-0059		F1562-0059_1_2 (1 of 4)

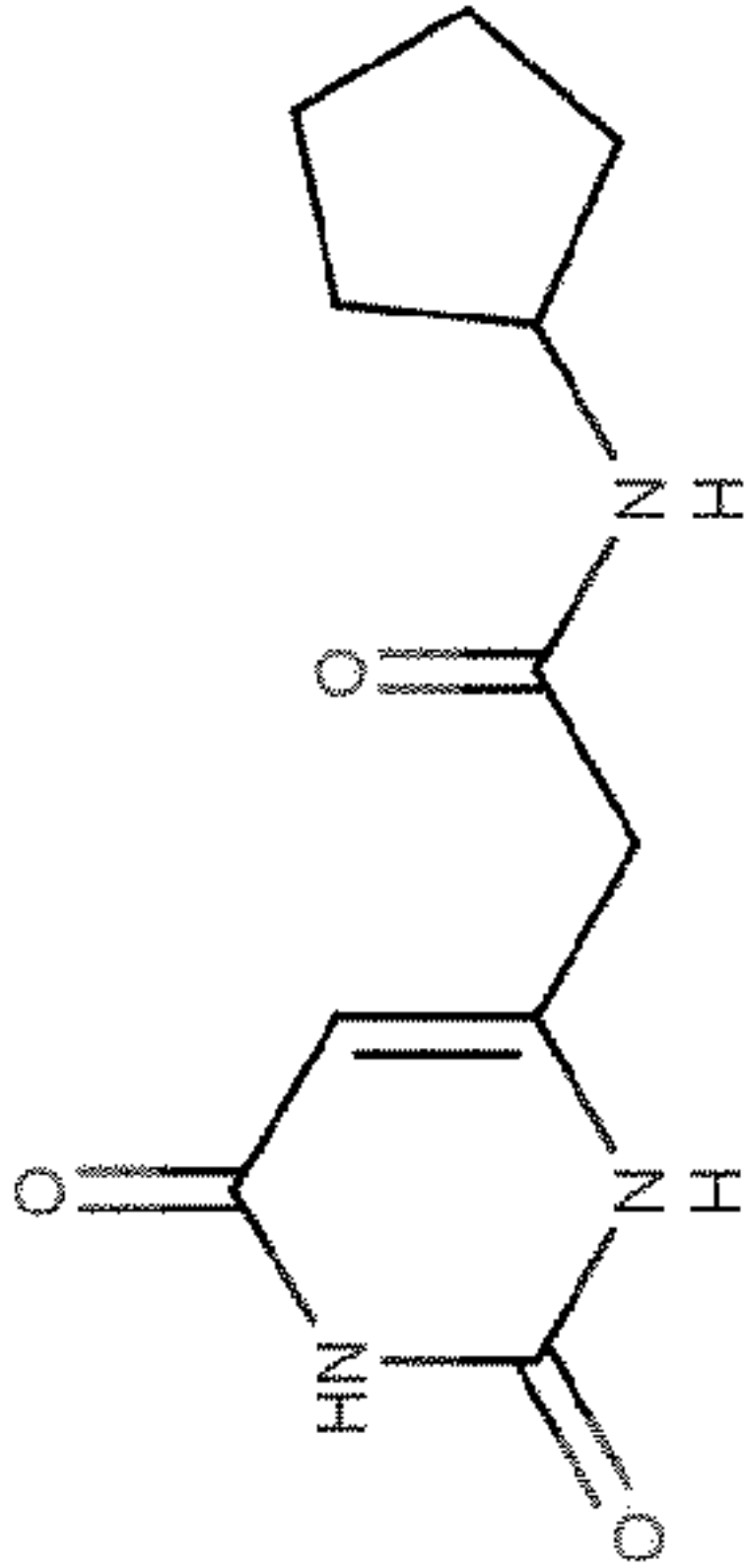
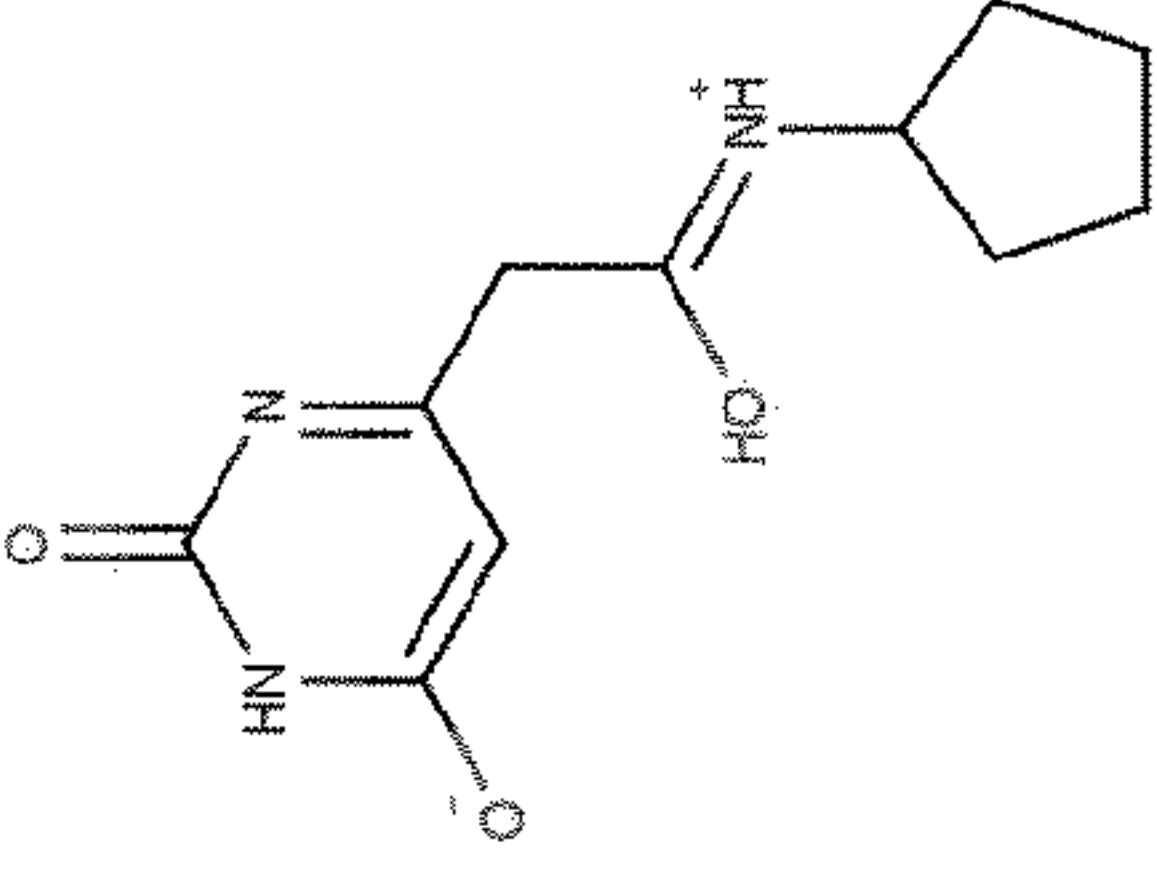
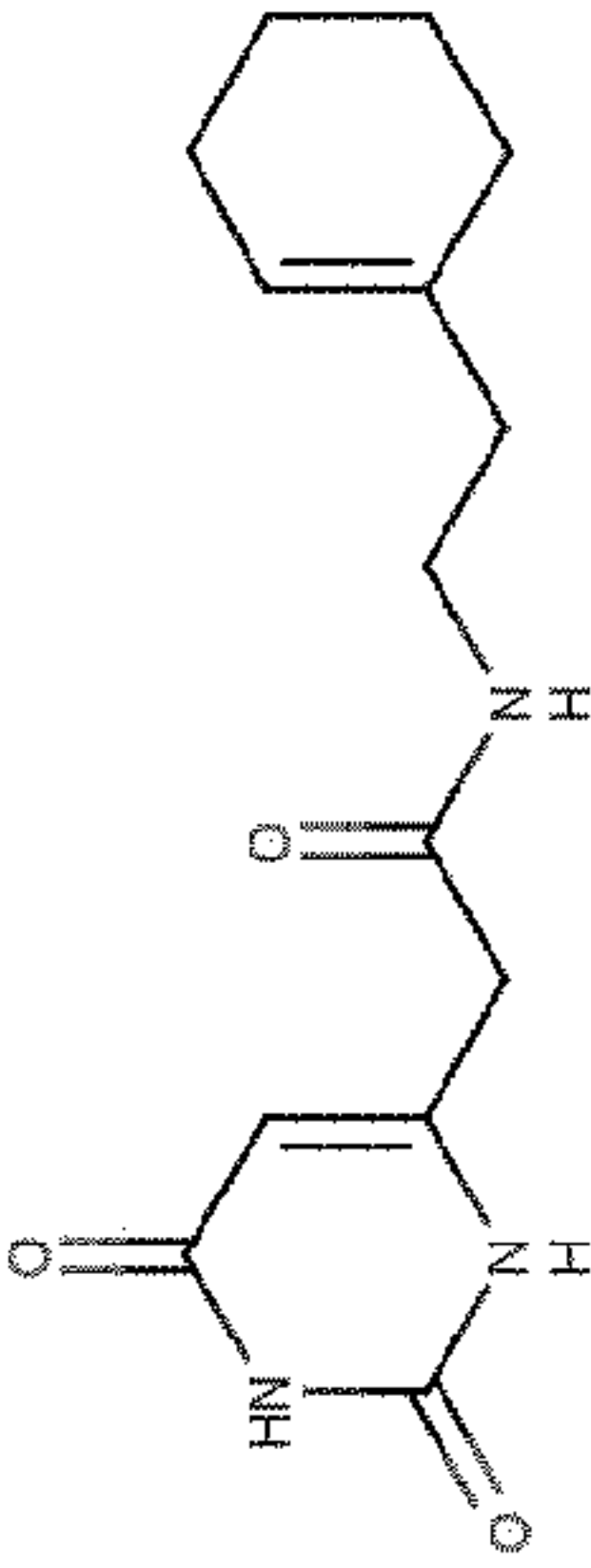
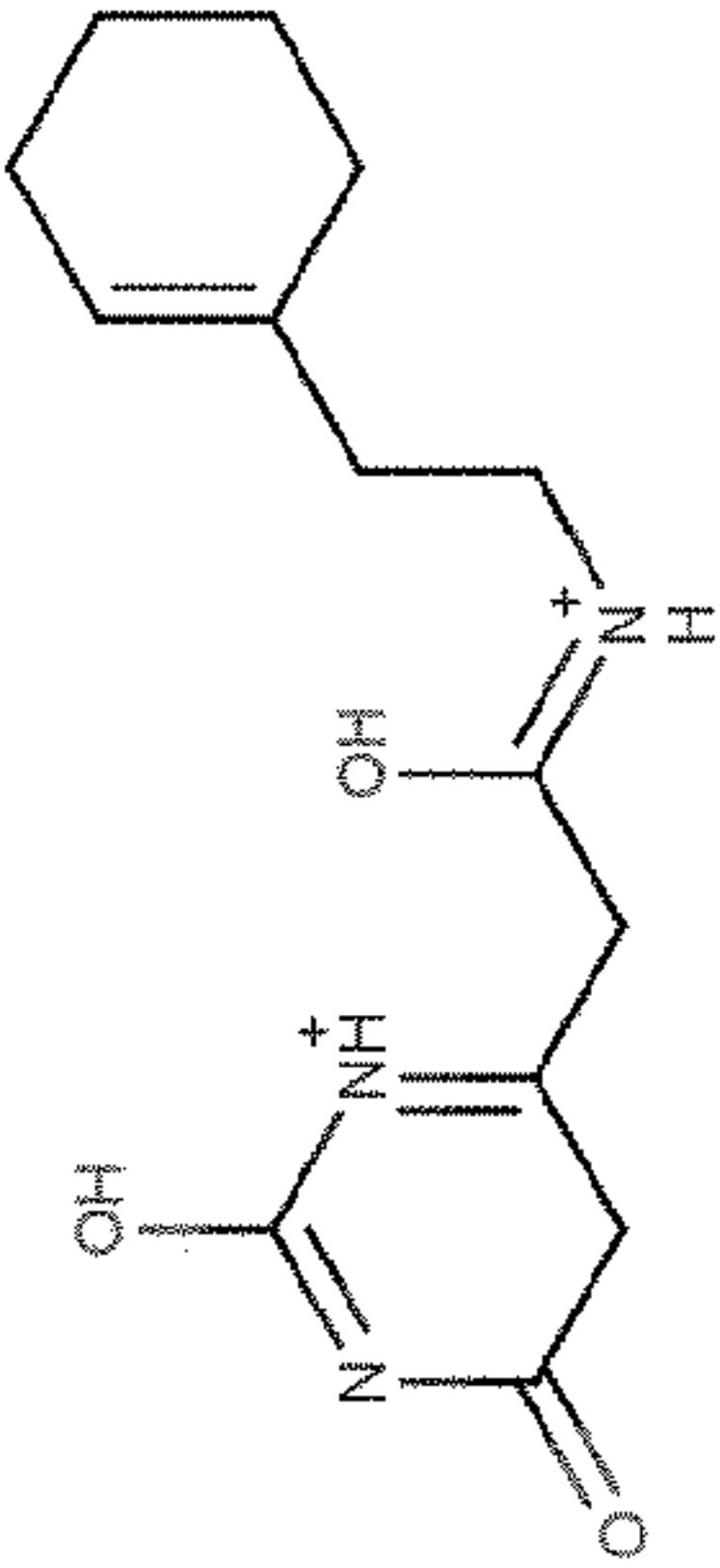
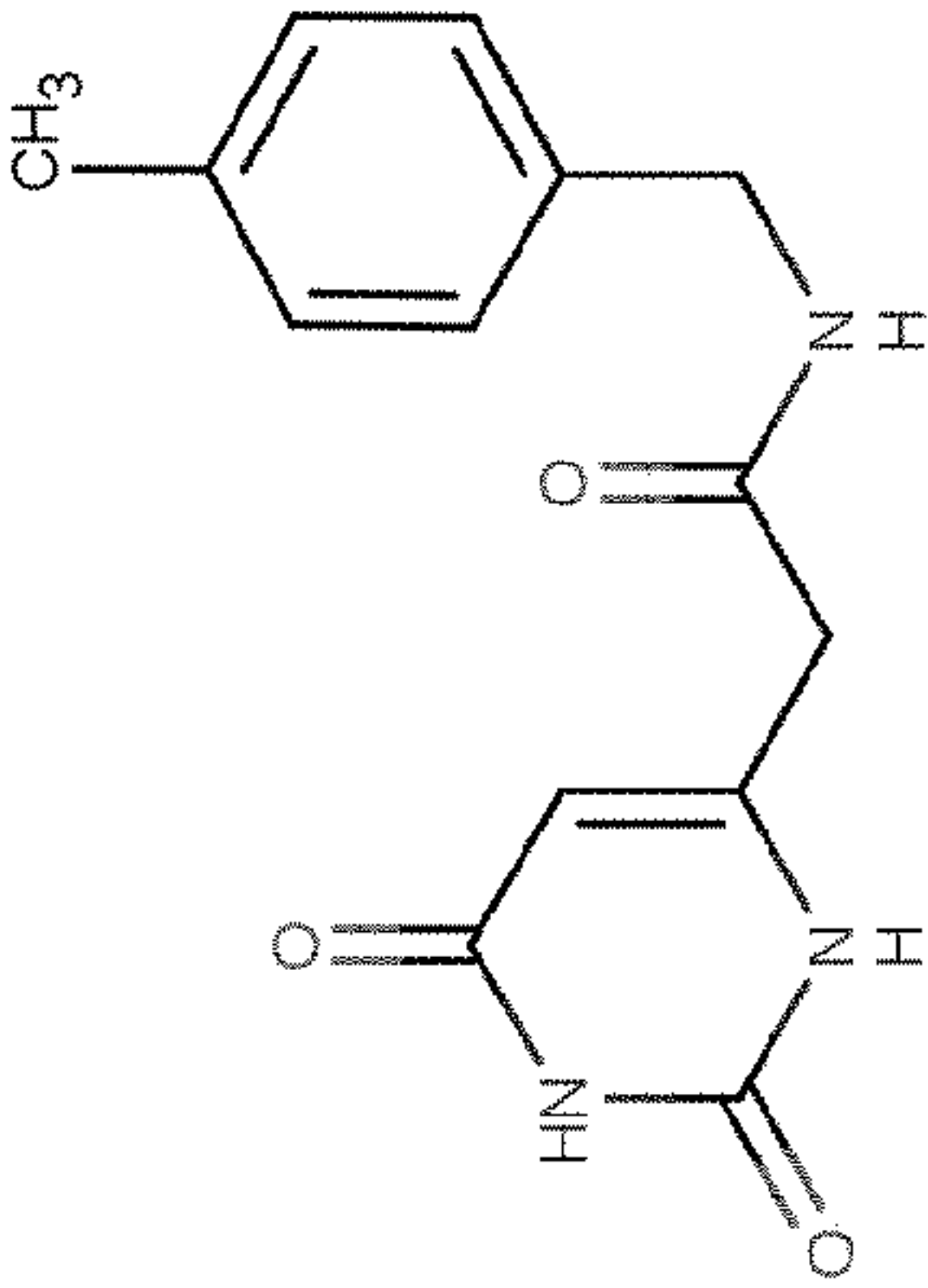
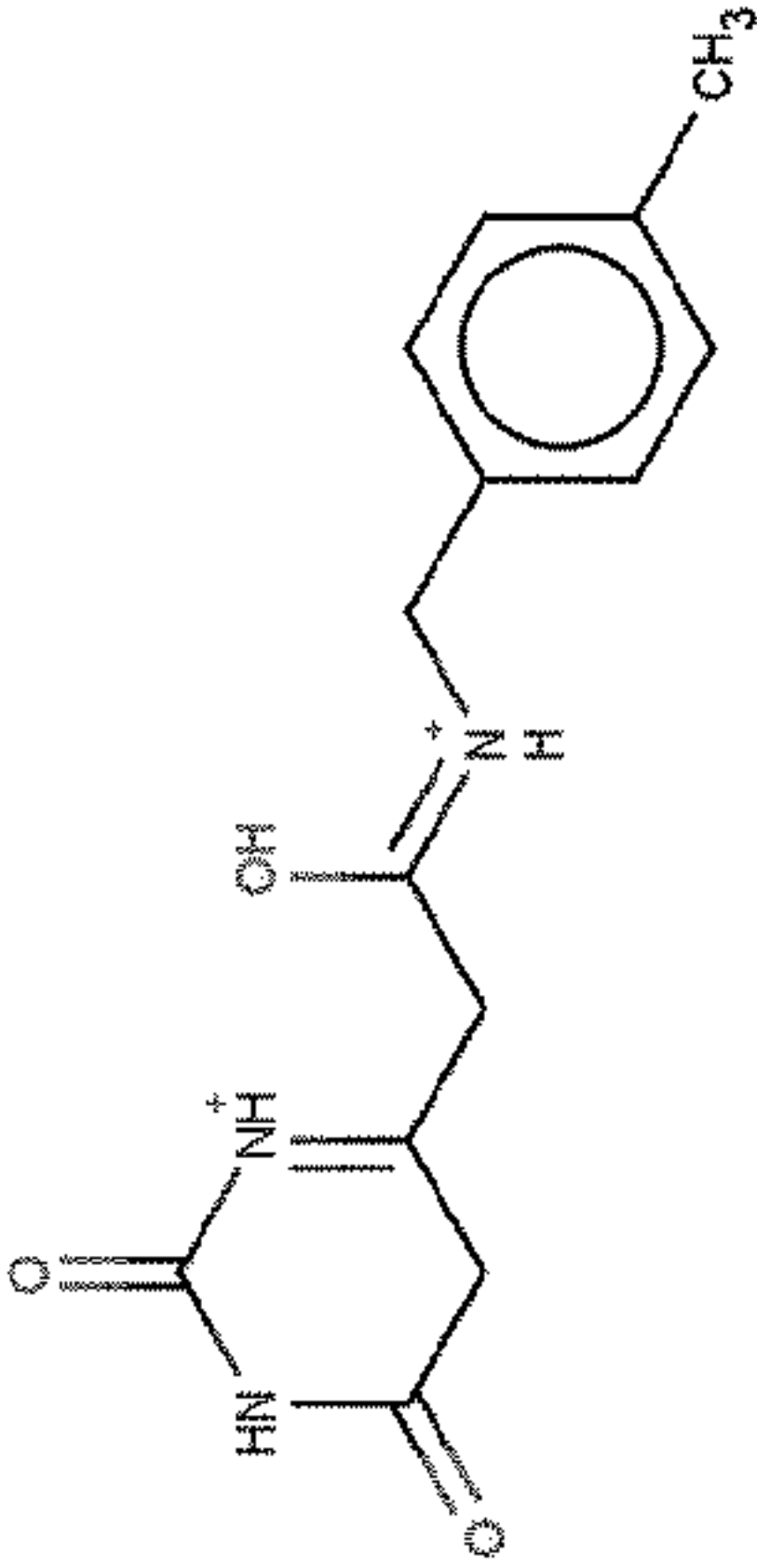
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 63		F1605-0391		F1605-0391_4_1 (1 of 37)
Compound 64		F1607-0384		F1607-0384_32_1 (1 of 66)
Compound 65		F1607-0387		F1607-0387_33_1 (1 of 33)

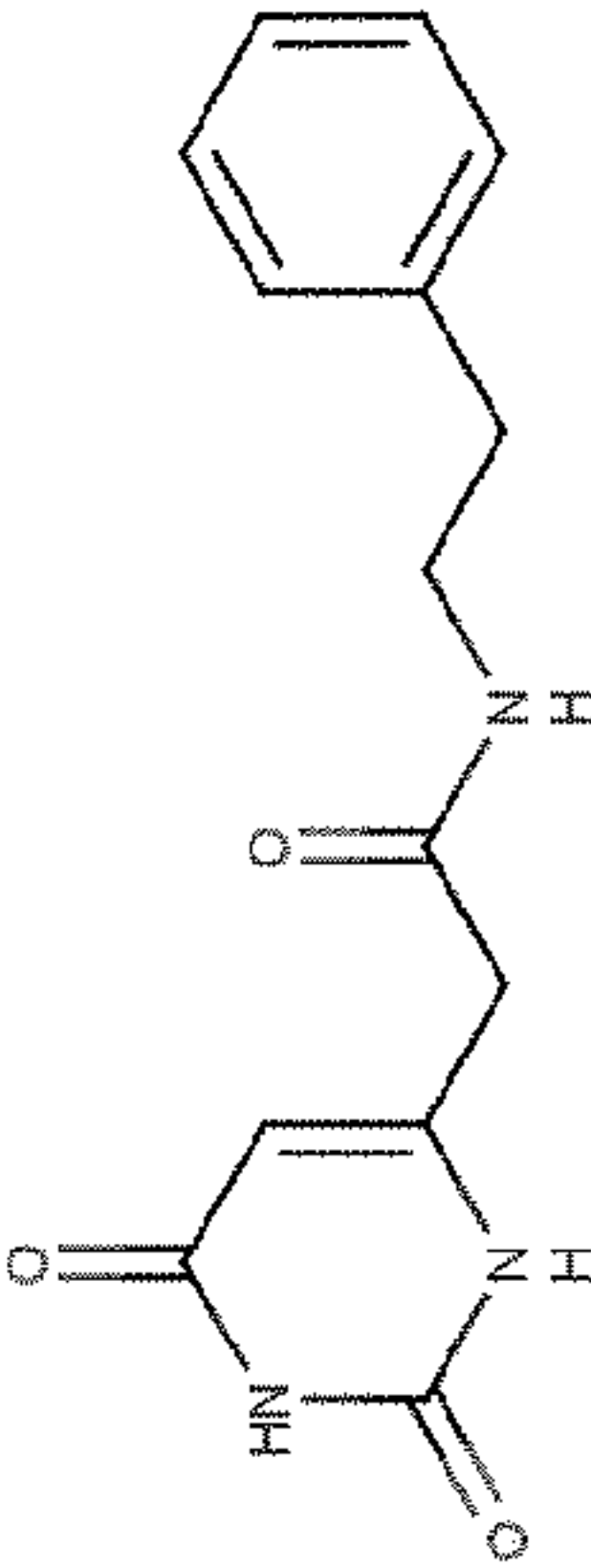
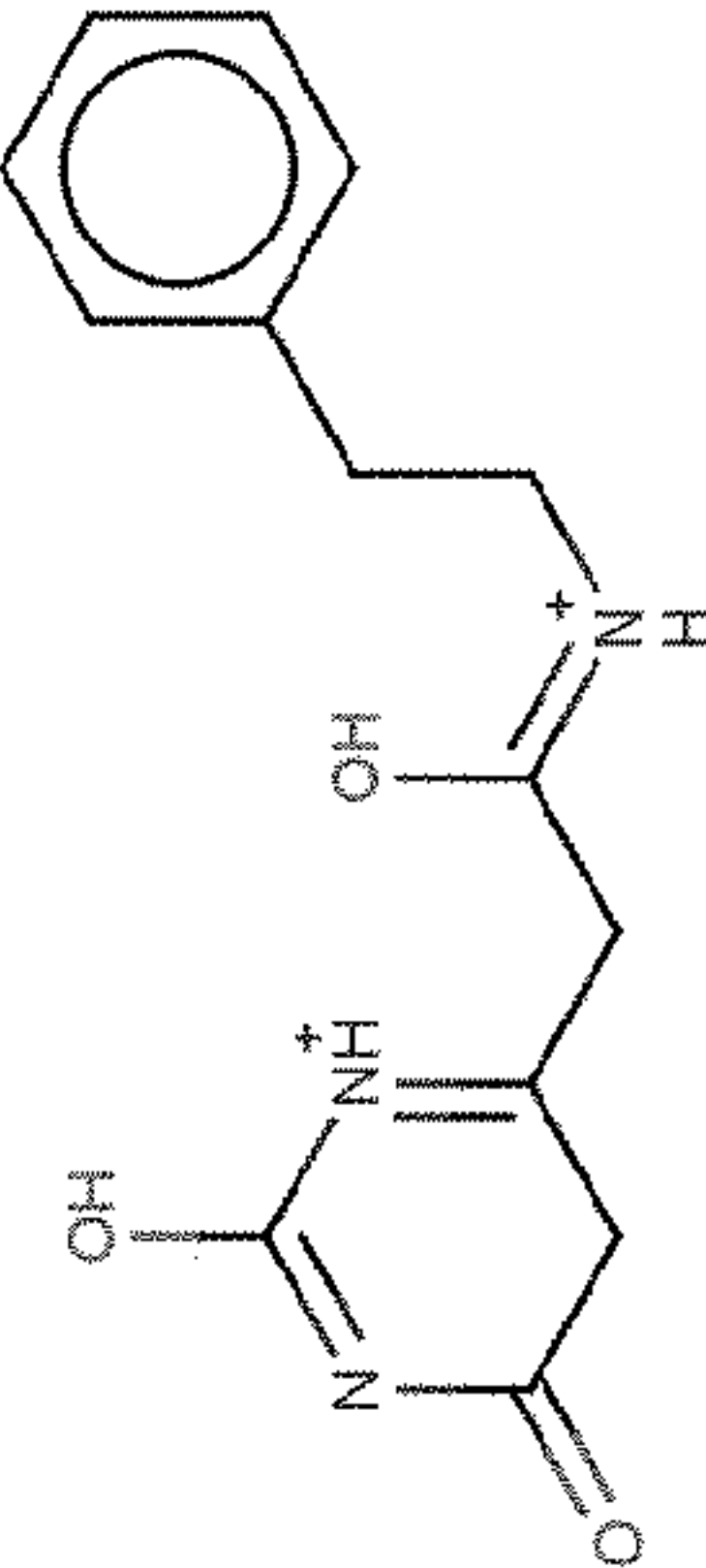
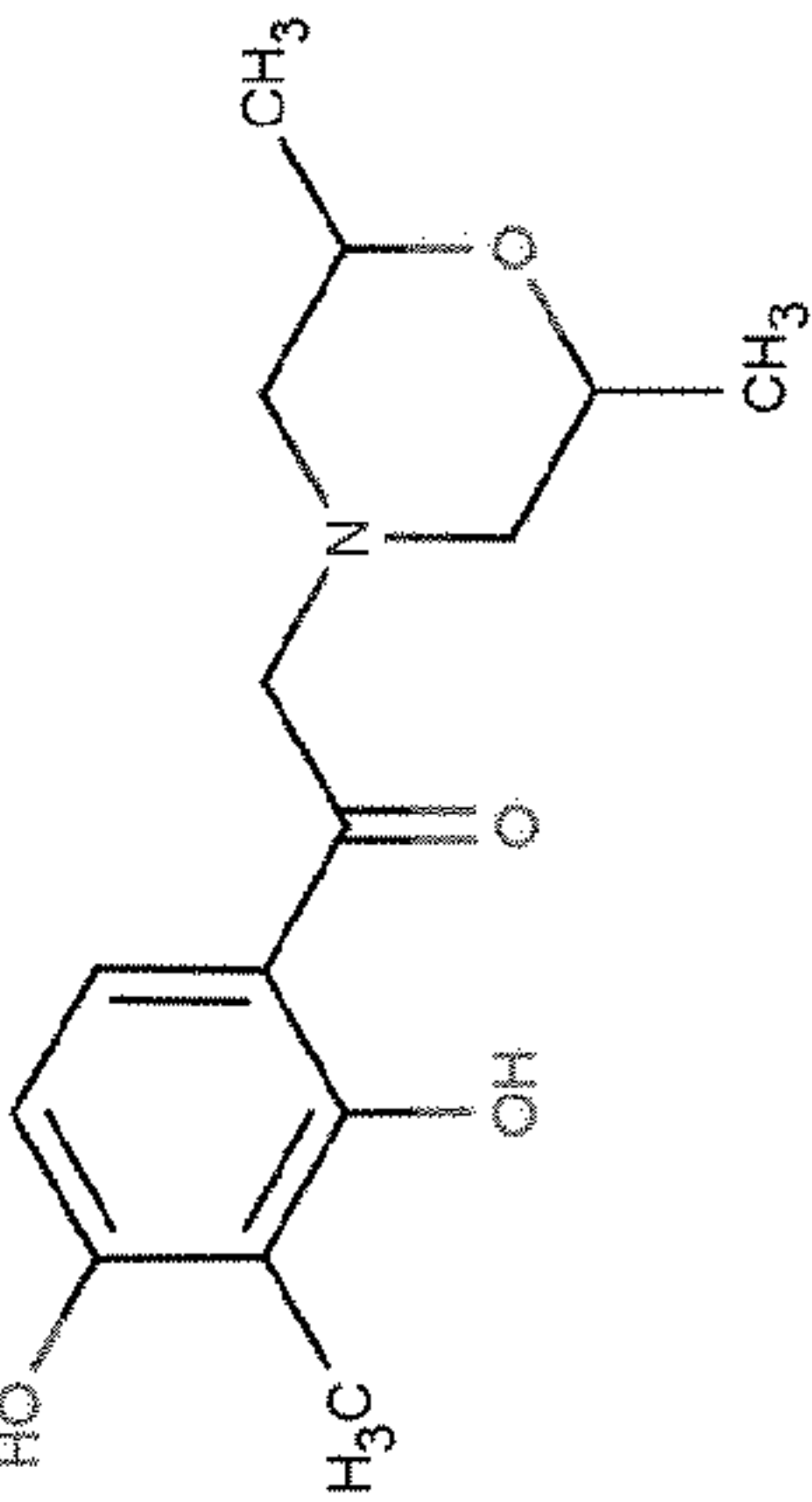
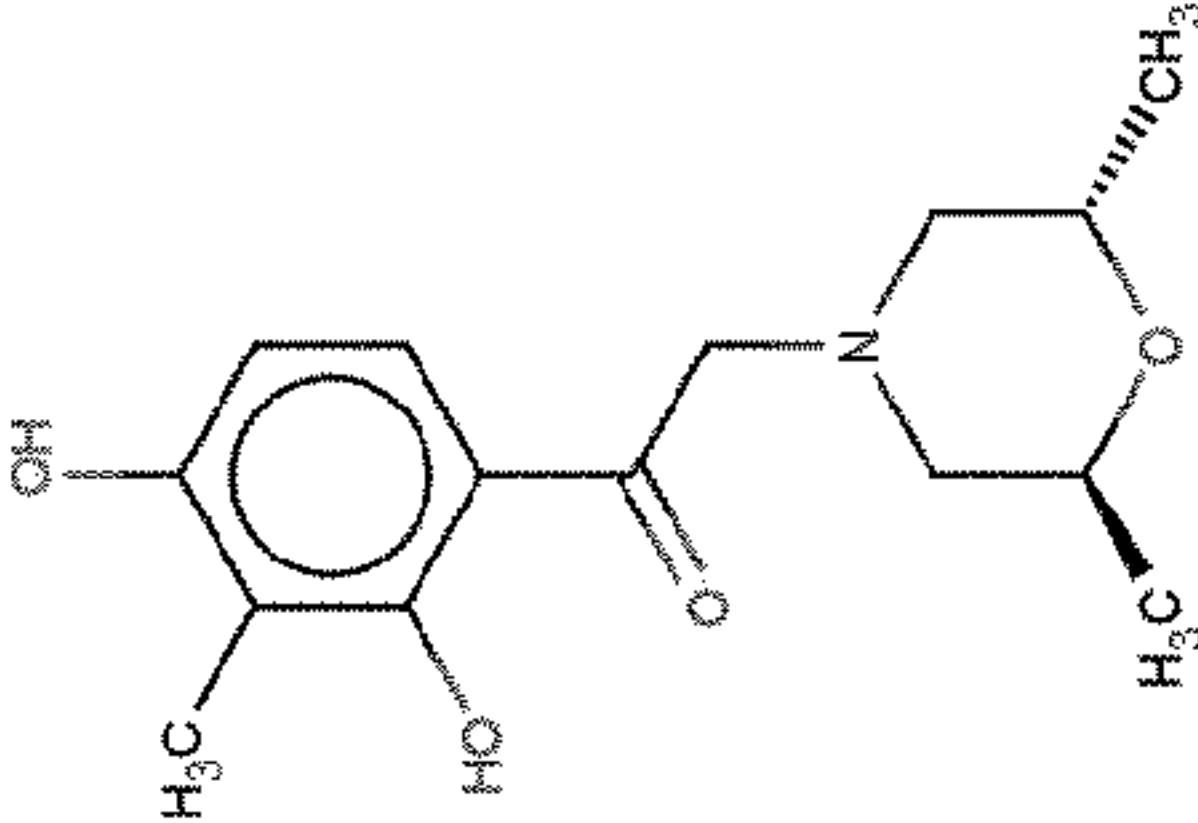
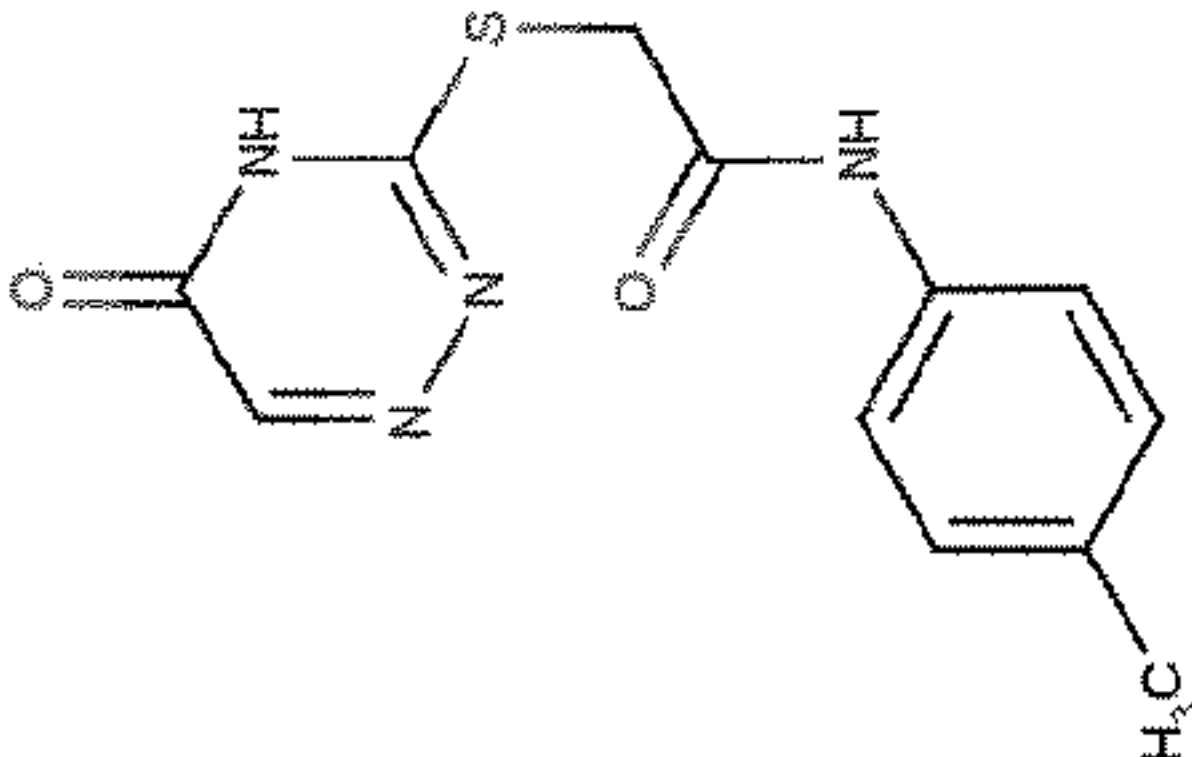
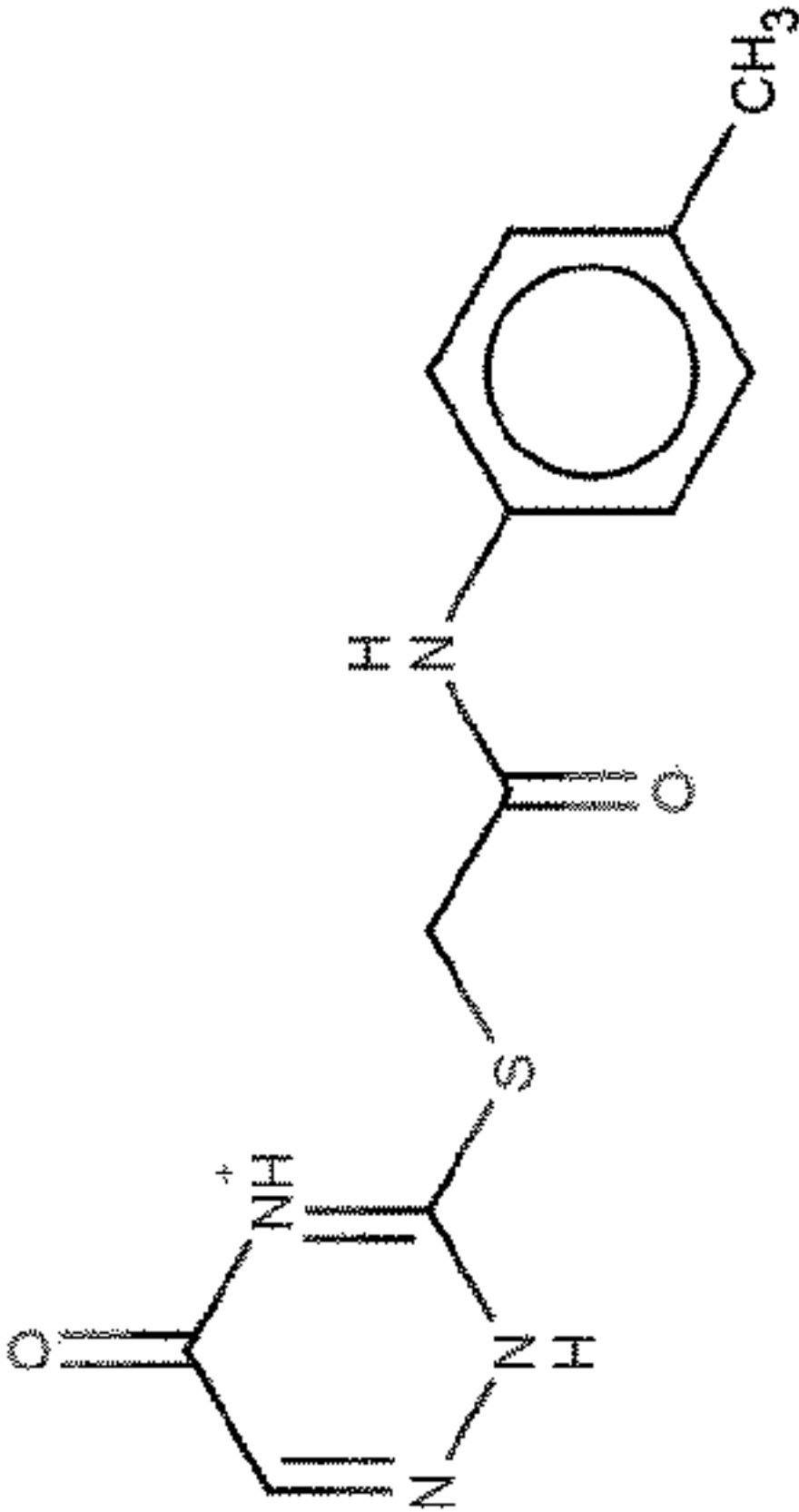
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 66		F1630-0018		F1630-0018_2 (1 of 4)
Compound 67		F1681-0025		F1681-0025_1_2 (1 of 4)
Compound 68		F1681-0068		F1681-0068_2 (1 of 4)

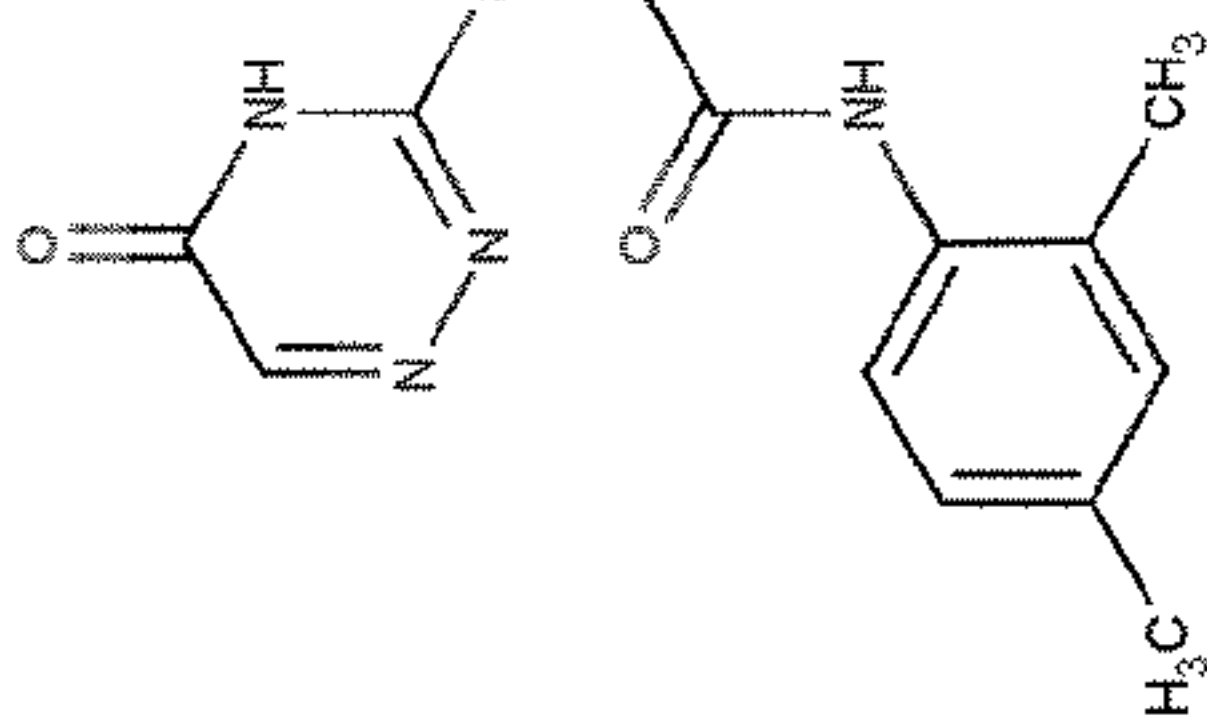
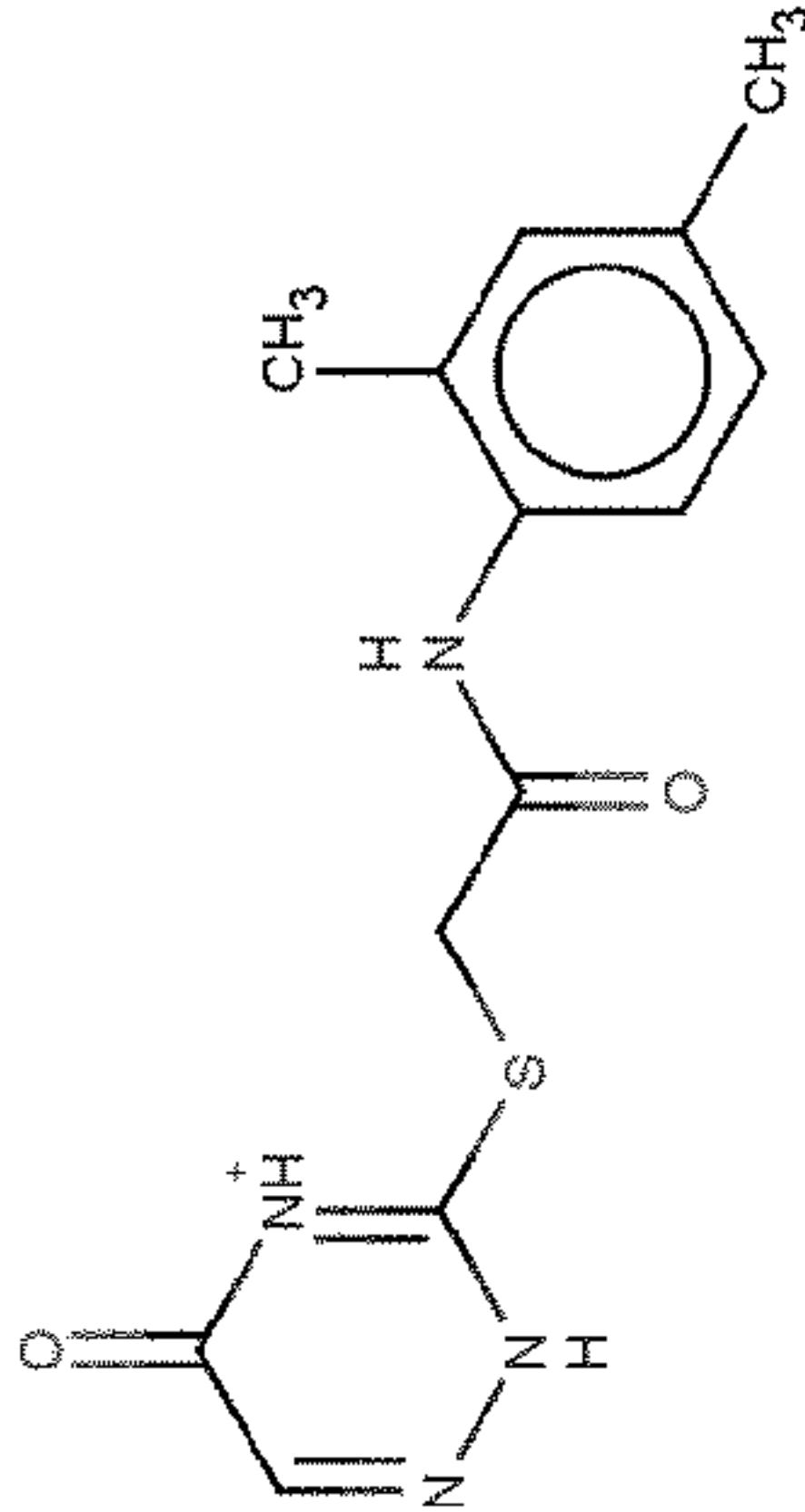
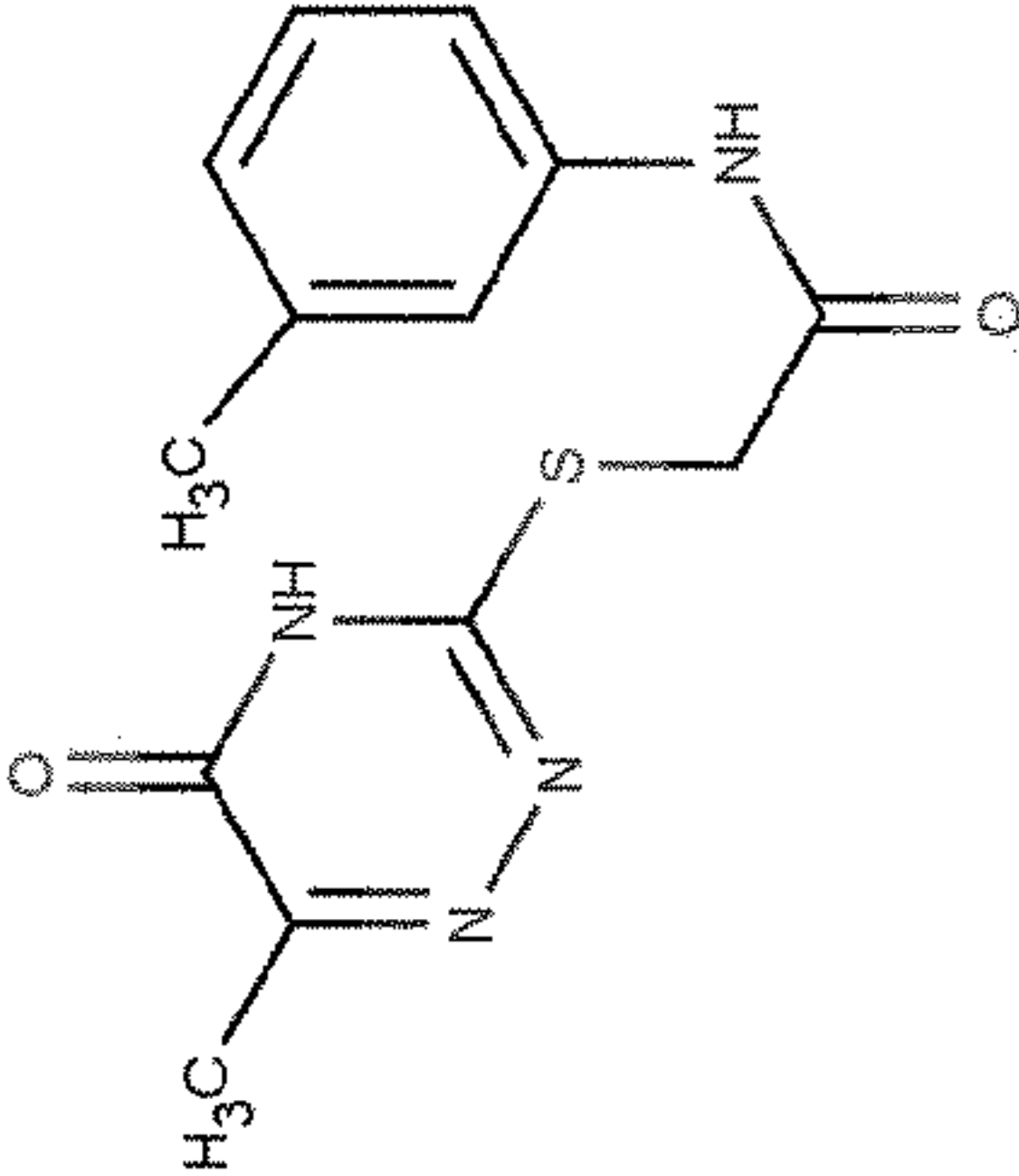
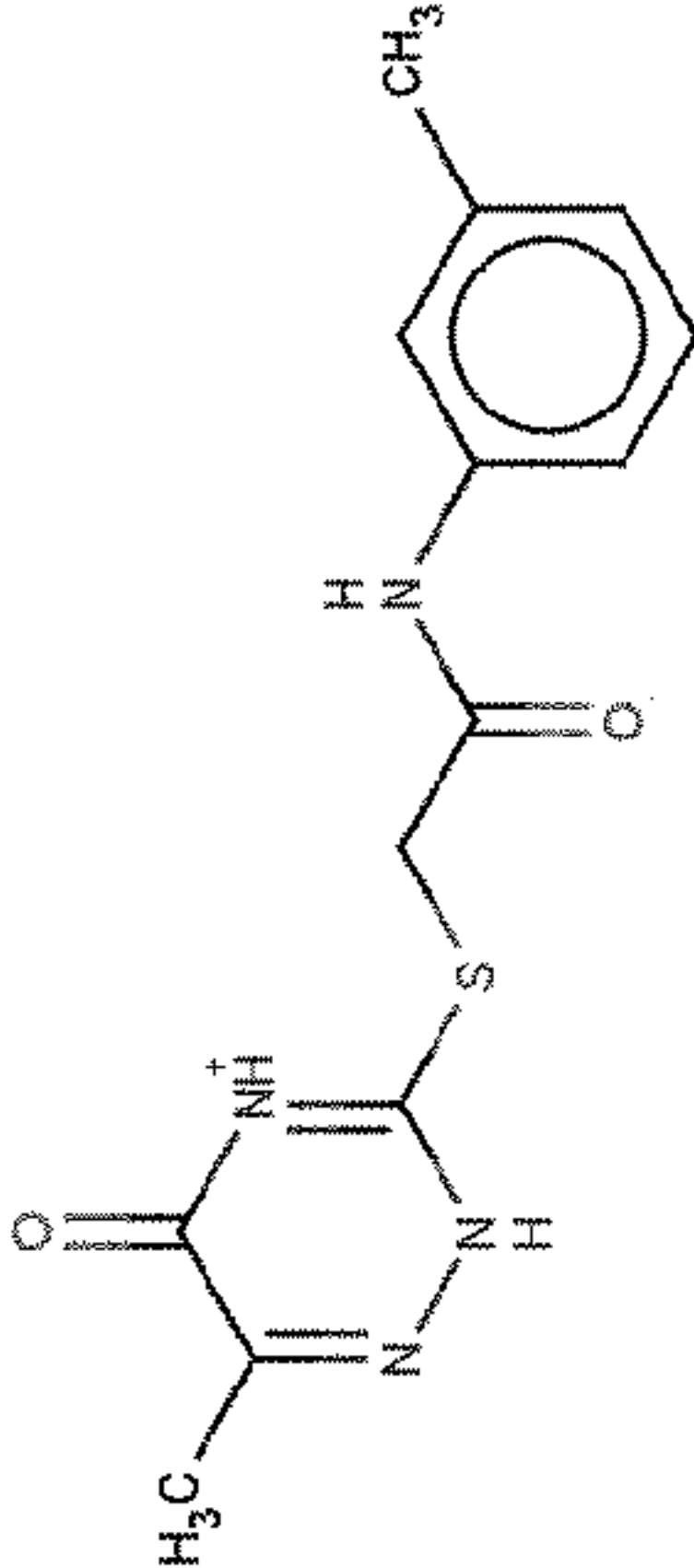
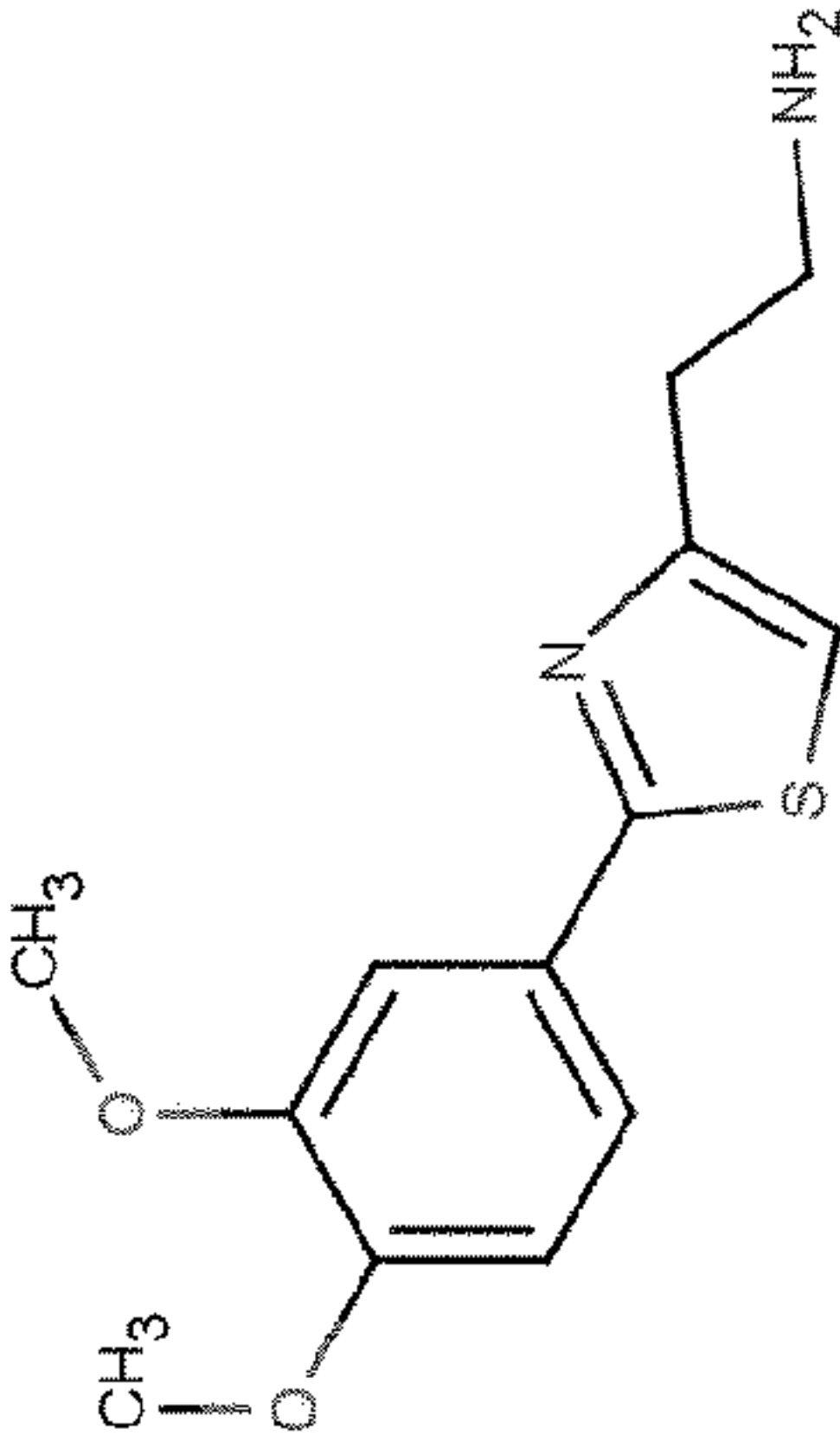
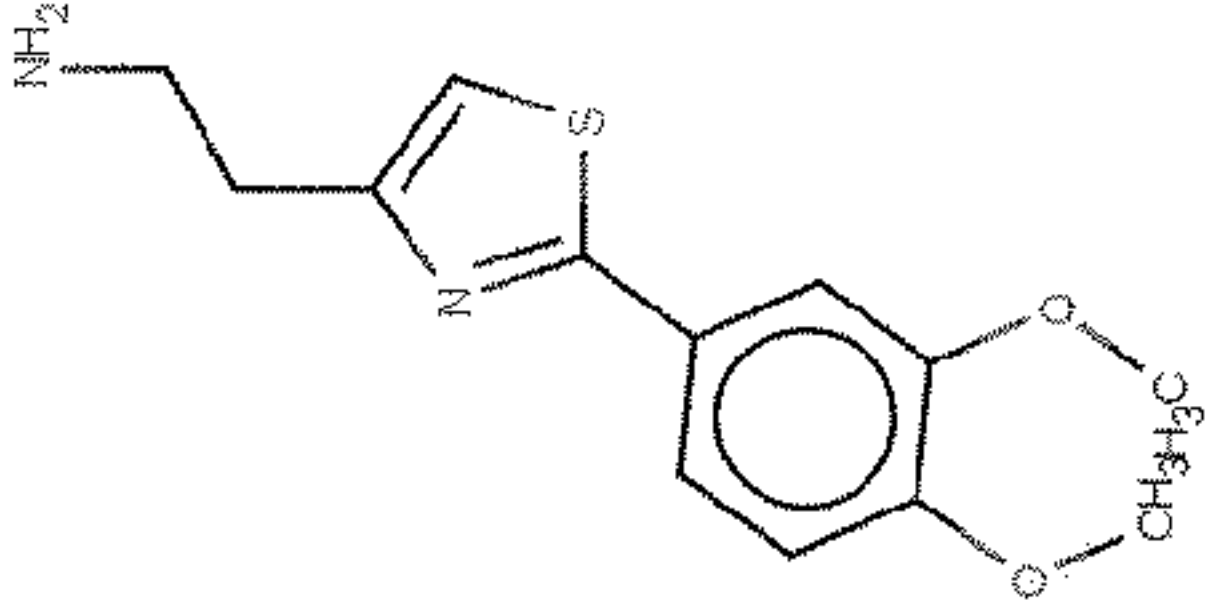
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 69		F1681-0091		F1681-0091_1 (1 of 4)
Compound 70		F1709-0008		F1709-0008_1_1 (1 of 4)
Compound 71		F1723-0297		F1723-0297_2_2 (1 of 8)

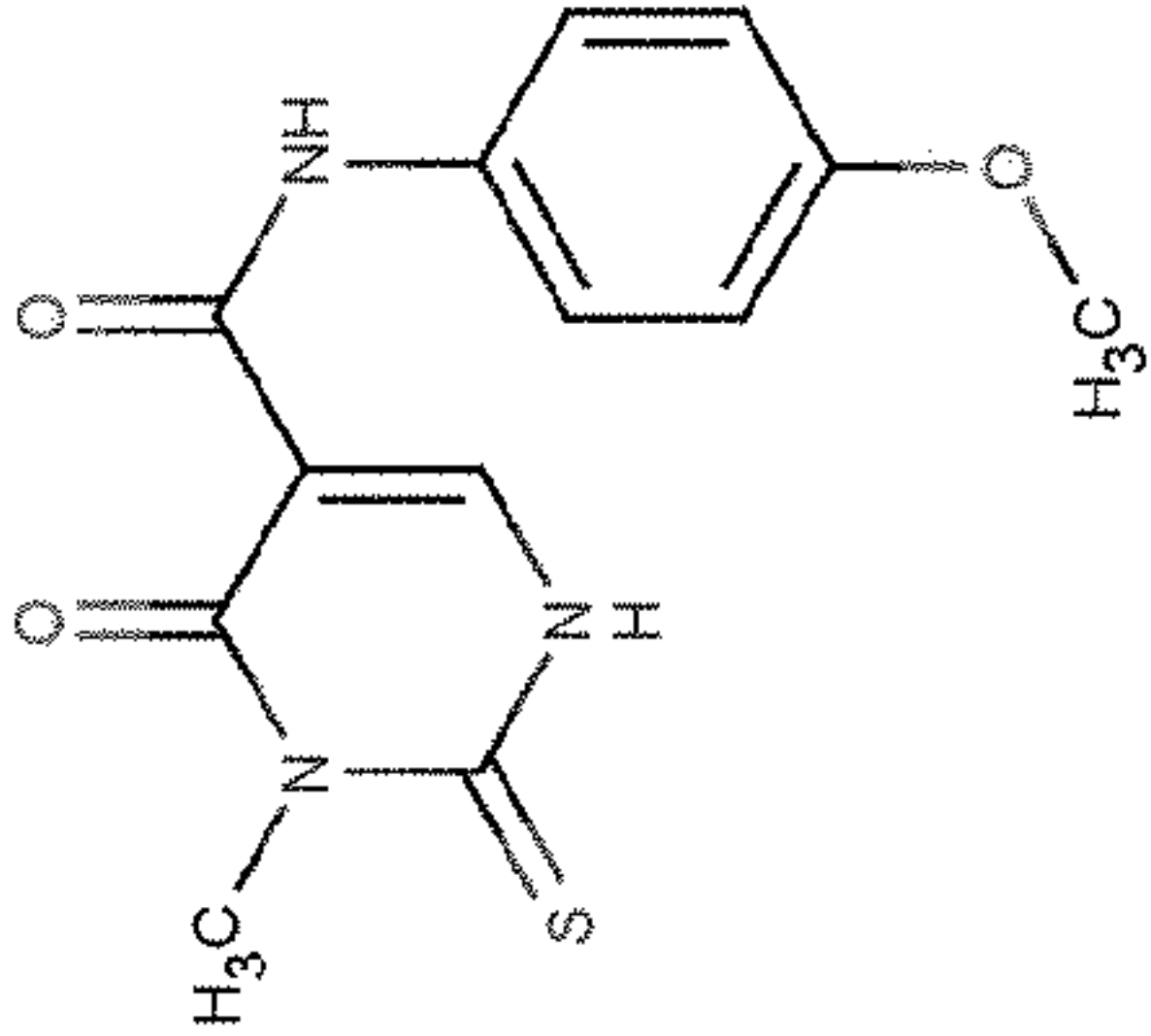
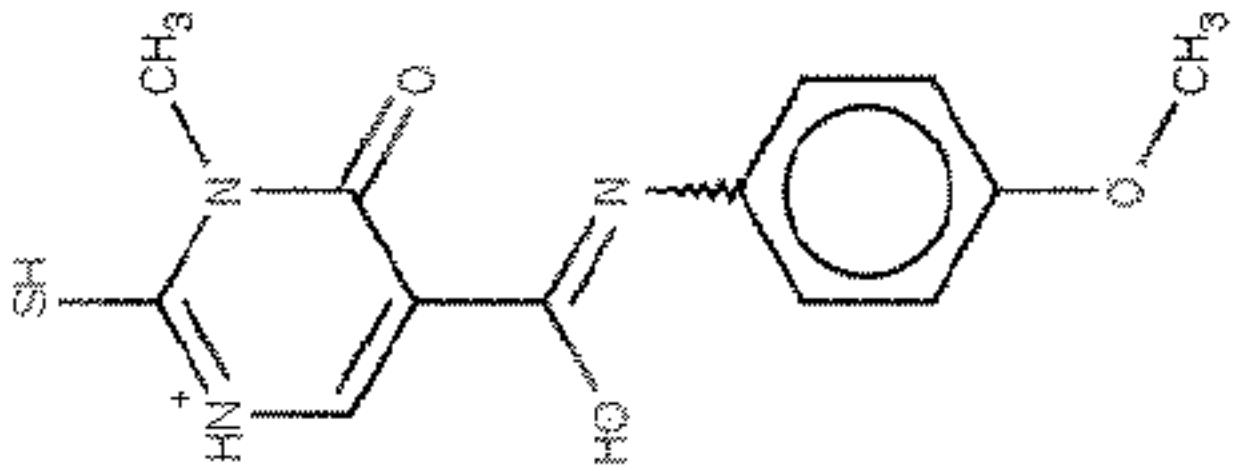
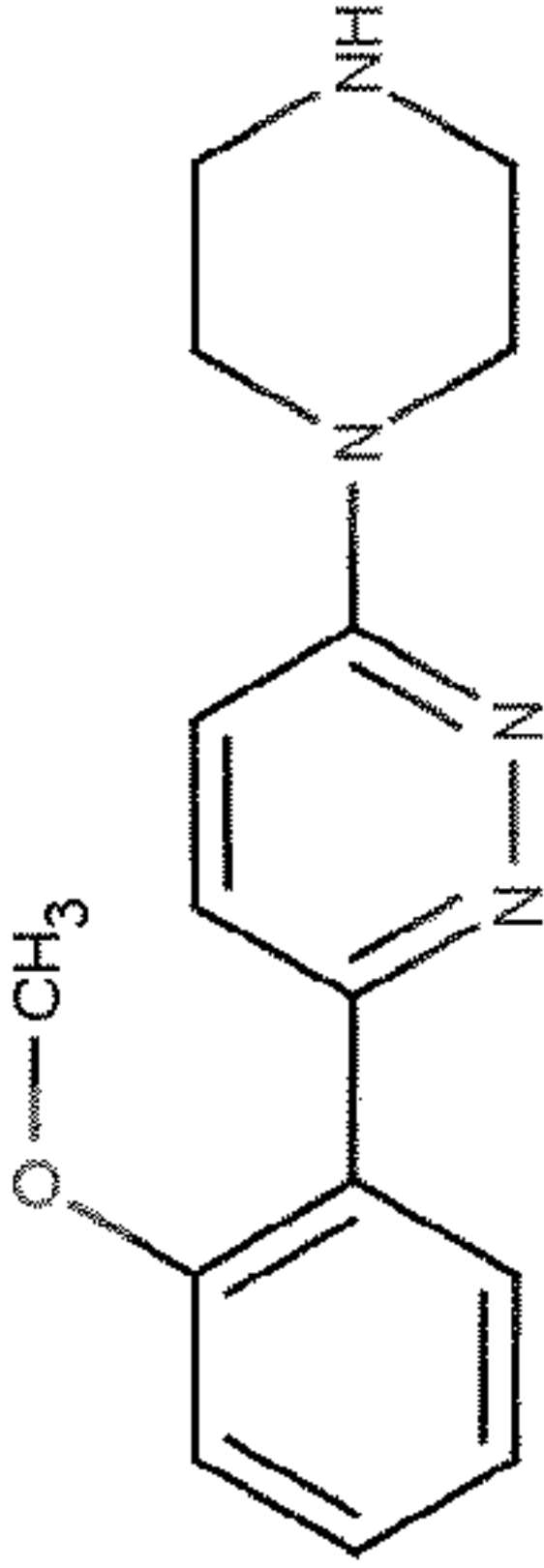
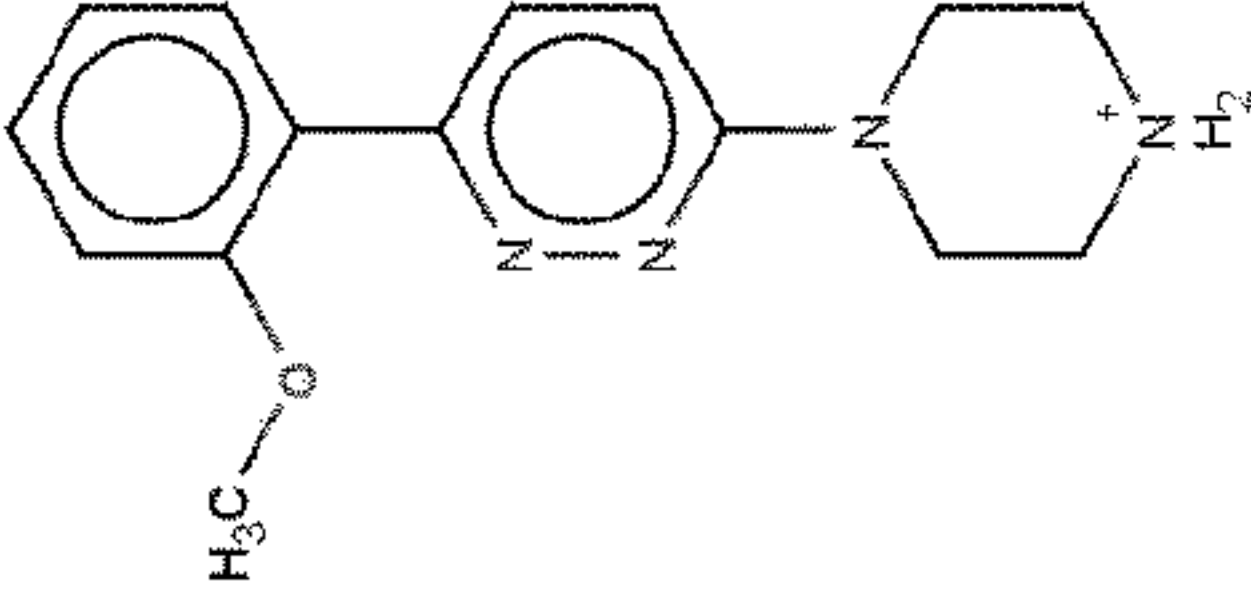
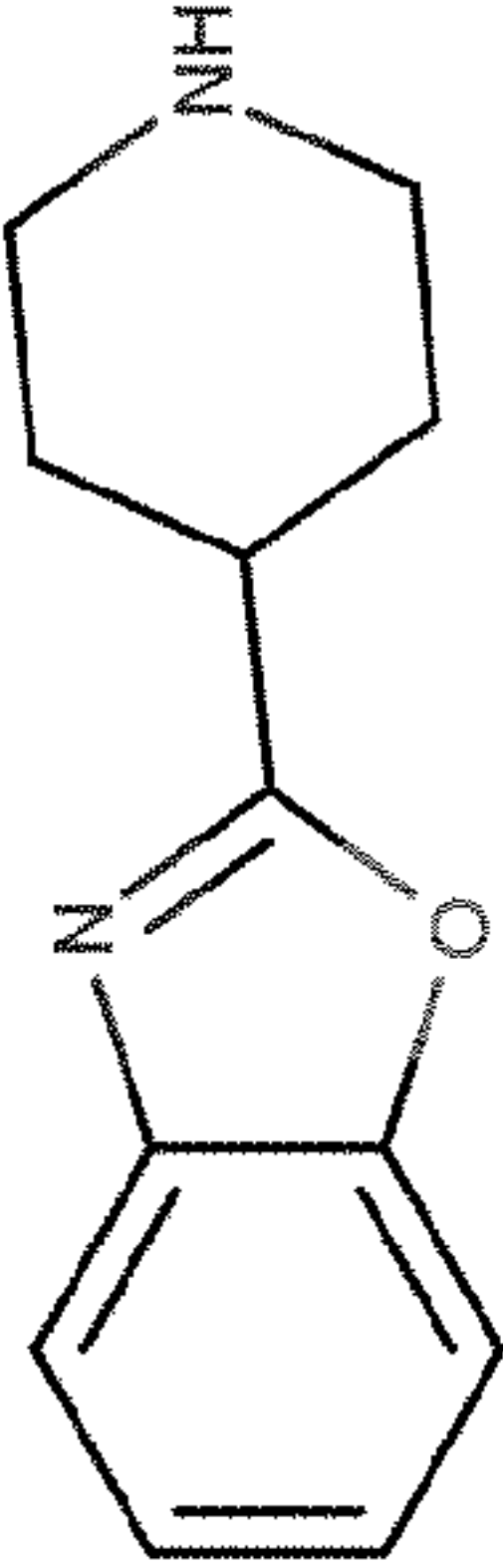
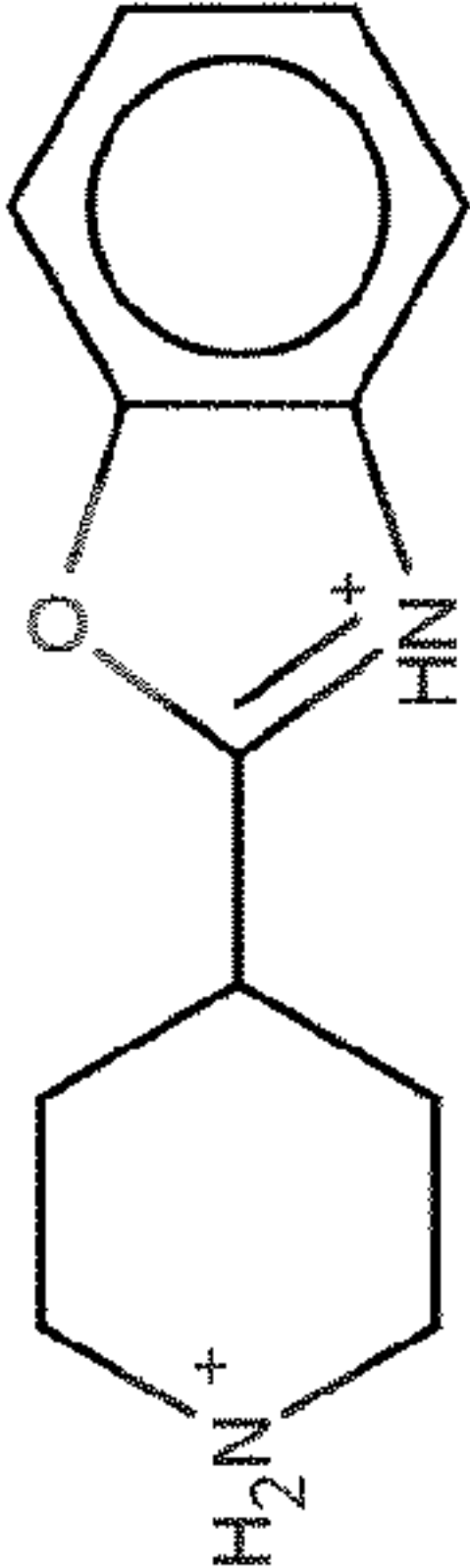
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 72		F1725-0012		F1725-0012_1_1 (1 of 4)
Compound 73		F1771-0001		F1771-0001_2 (1 of 4)
Compound 74		F1771-0003		F1771-0003_1_2 (1 of 4)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 75		F1771-0007		F1771-0007_1_8 (1 of 16)
Compound 76		F1811-0028		F1811-0028_57_1 (1 of 139)
Compound 77		F1811-0068		F1811-0068_142_1 (1 of 211)

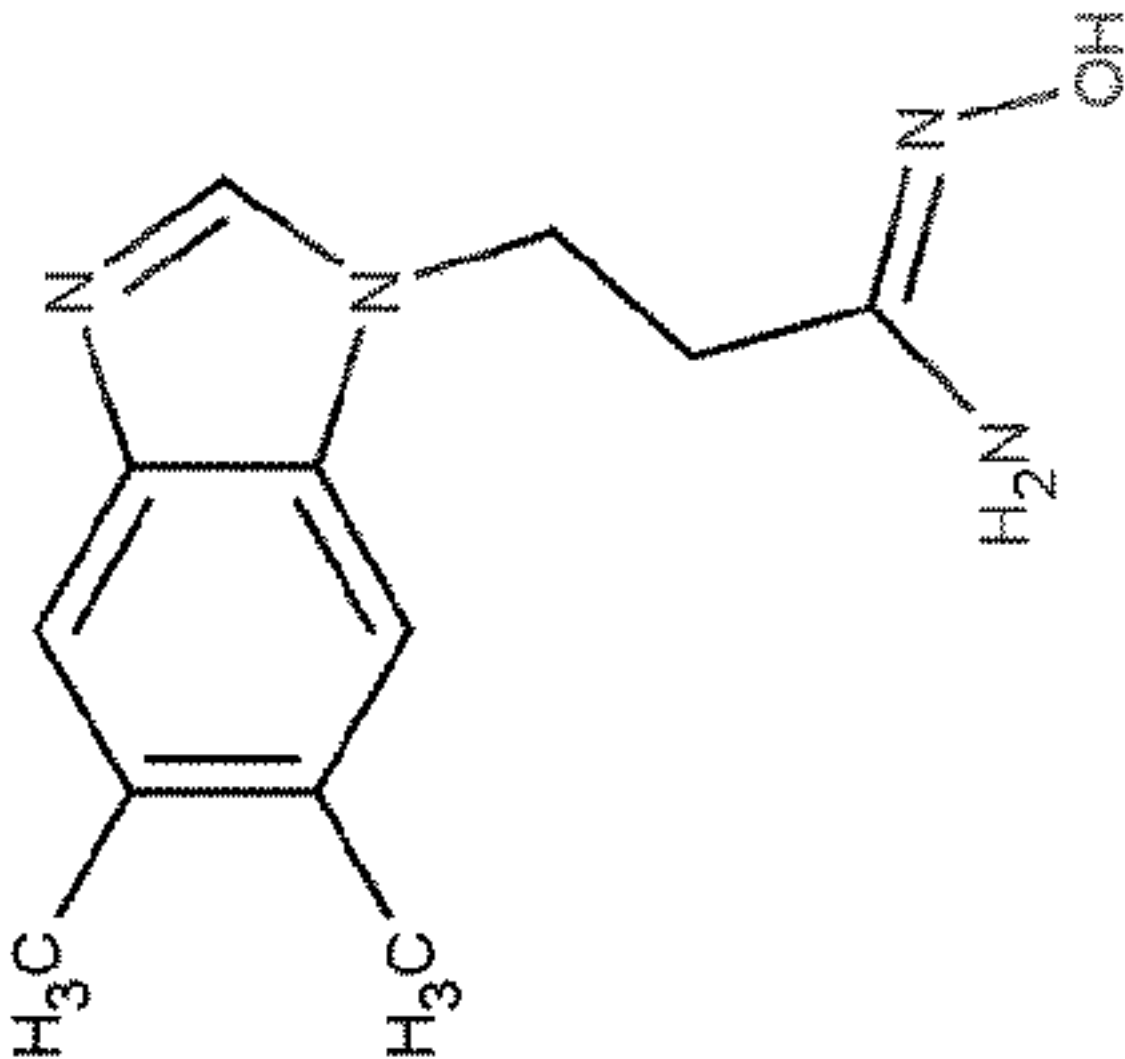
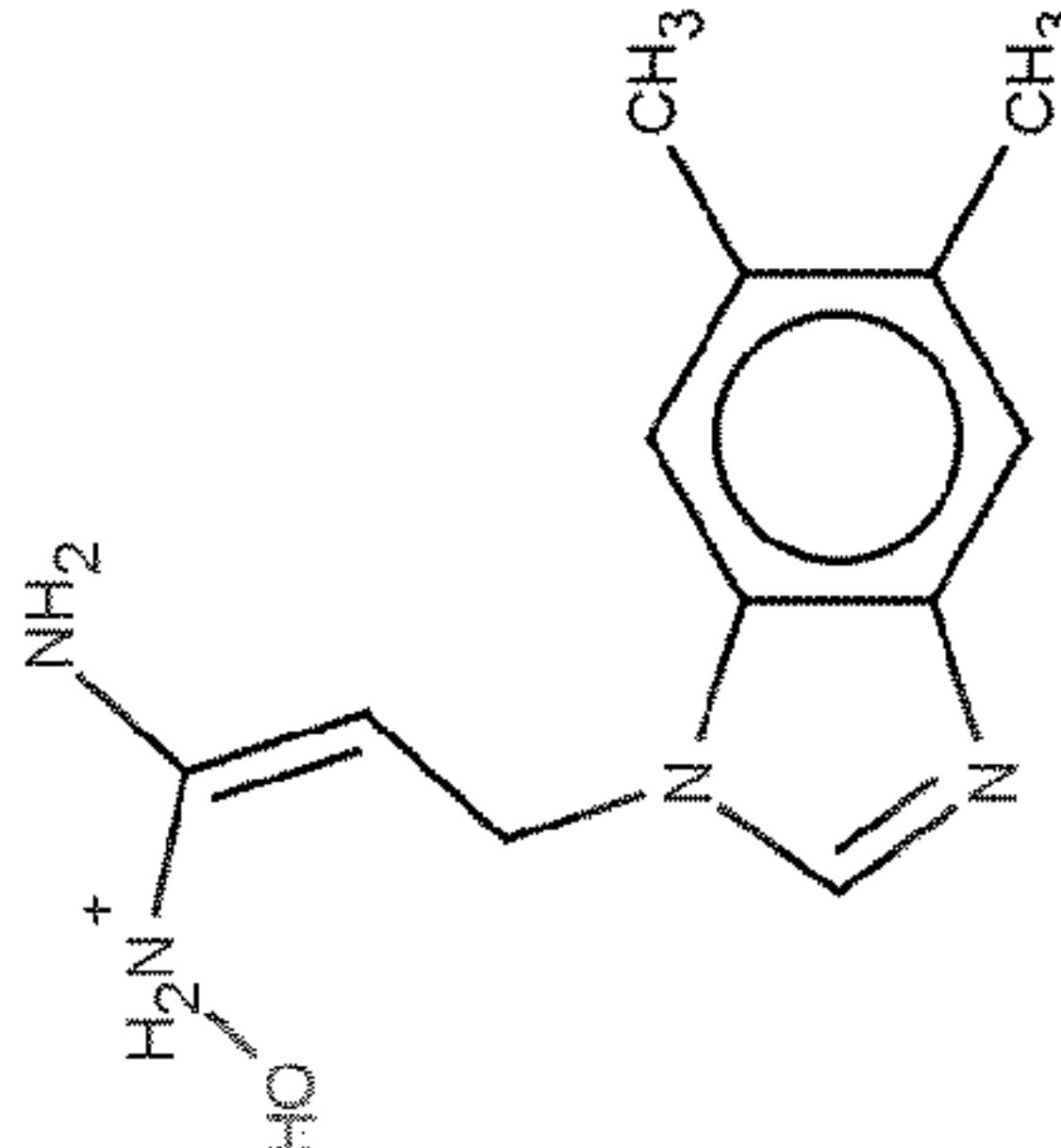
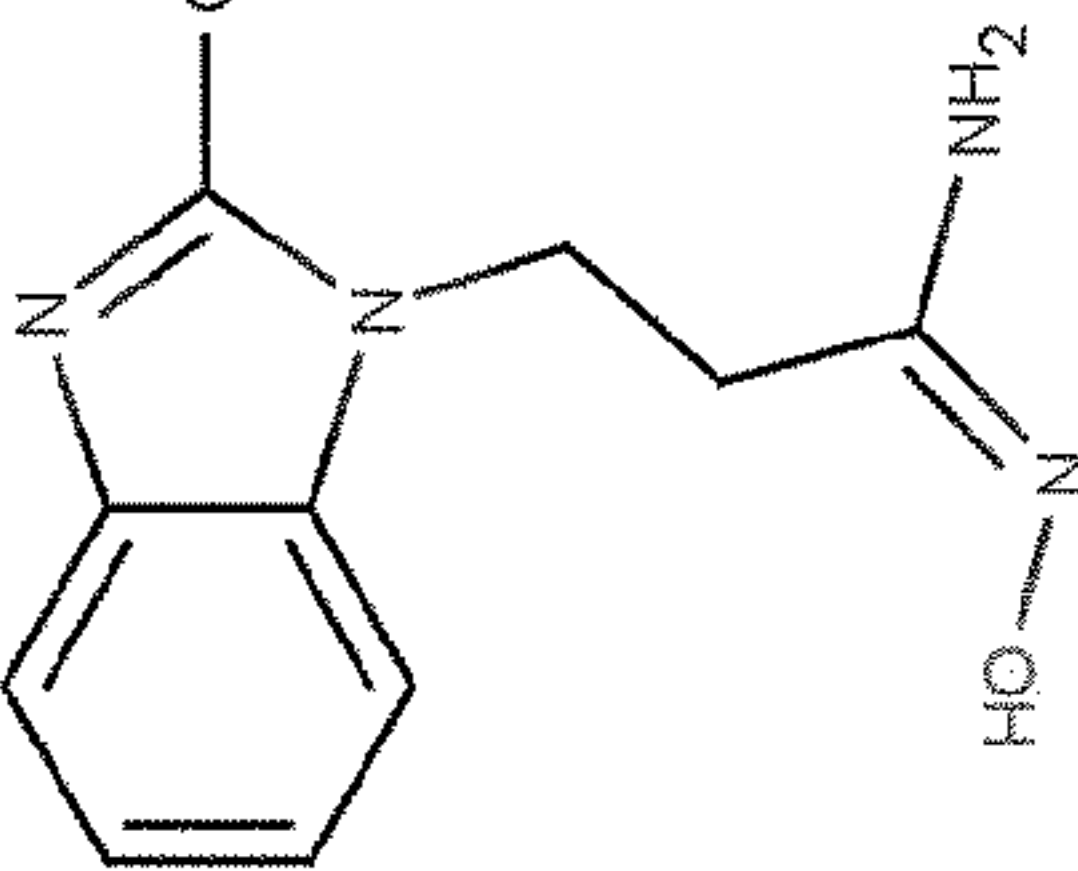
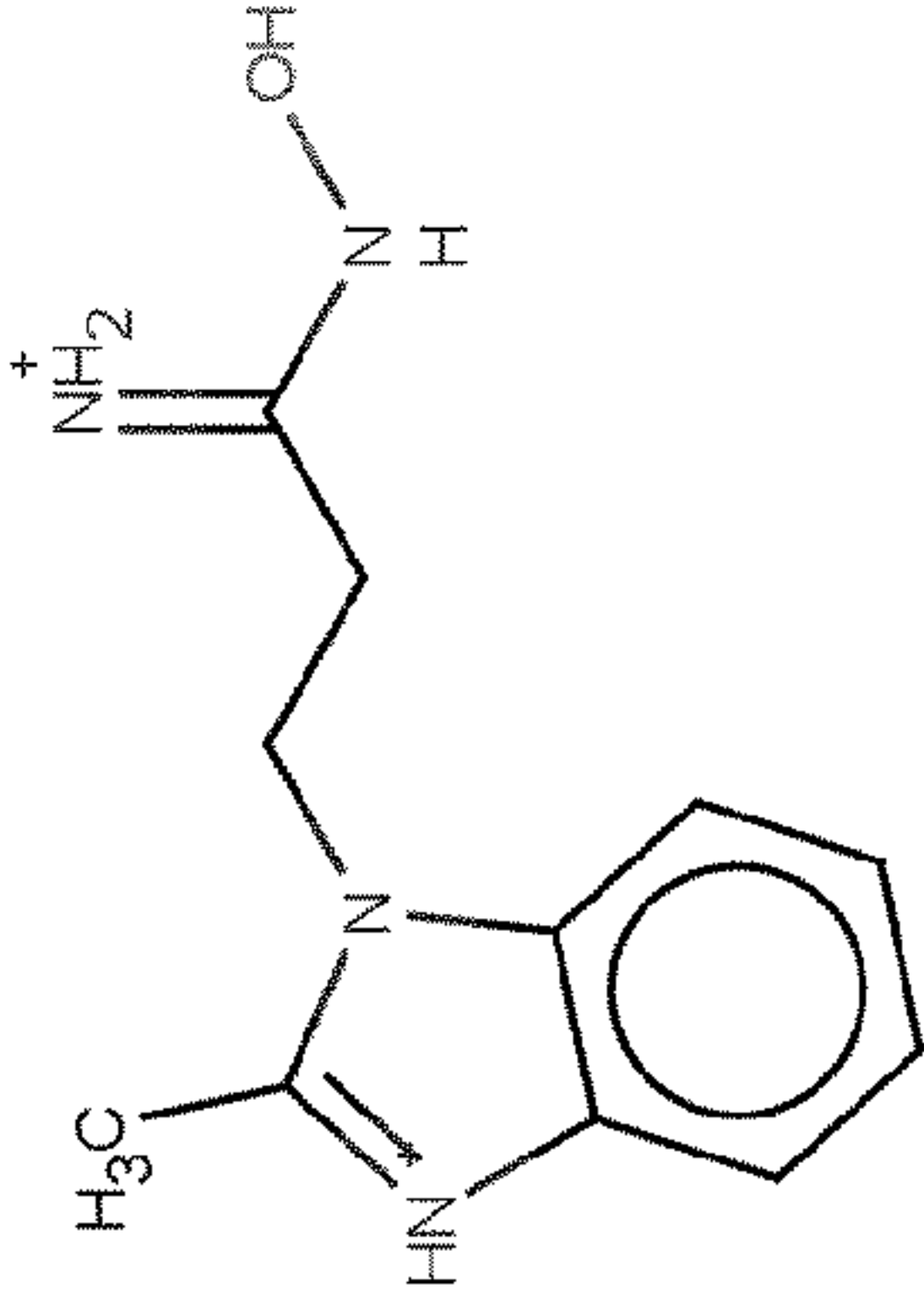
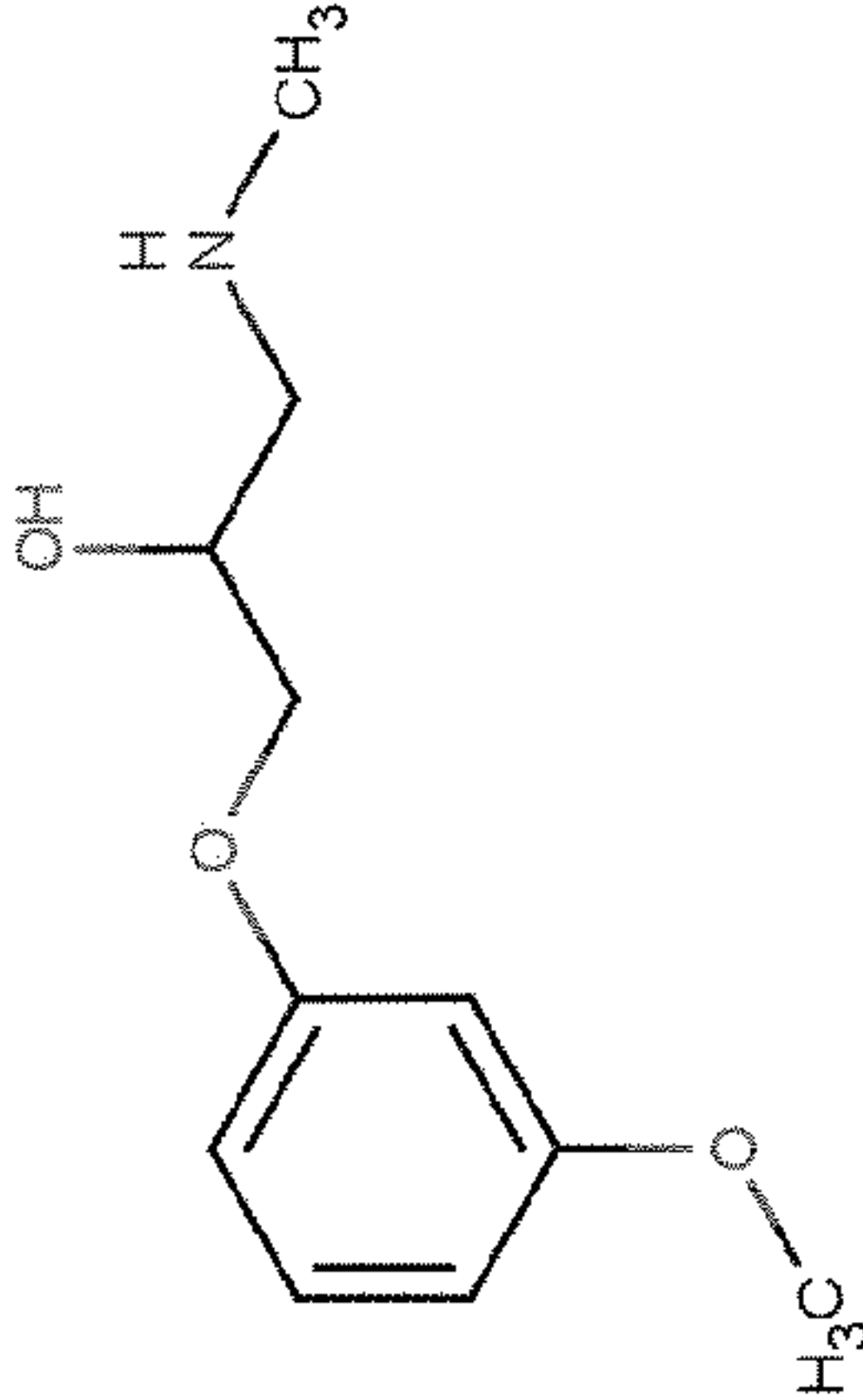
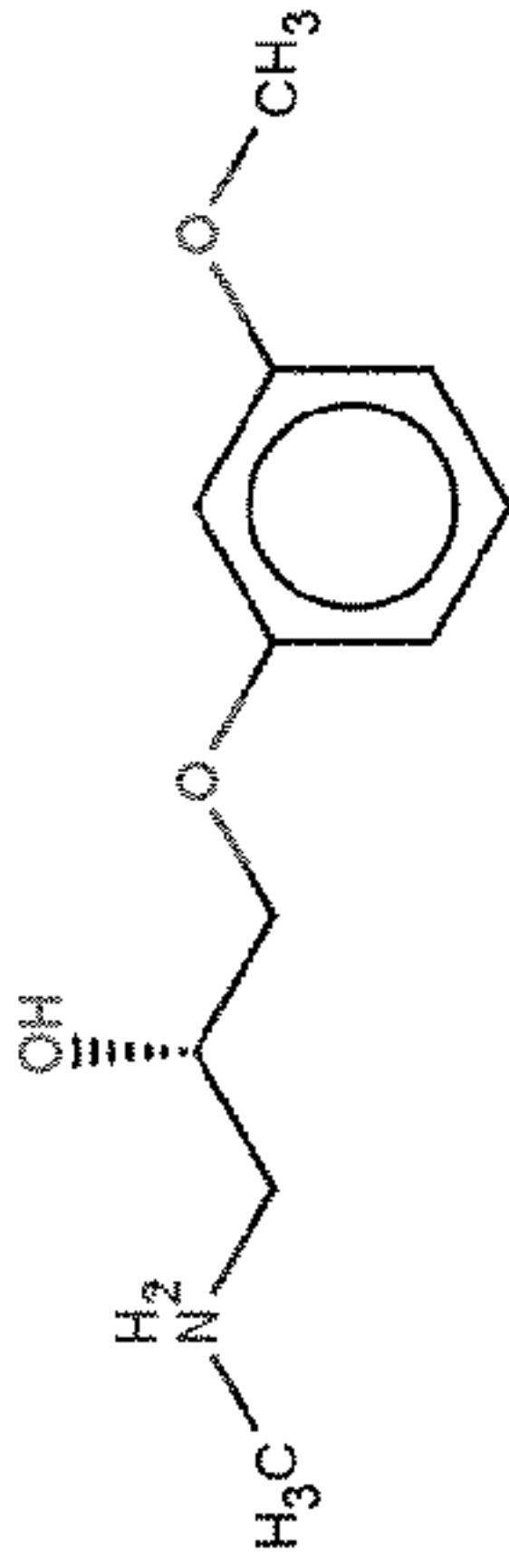
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 78		F1811-0098		F1811-0098_148_1 (1 of 238)
Compound 79		F1811-0101		F1811-0101_142_1 (1 of 203)
Compound 80		F1811-0112		F1811-0112_137_1 (1 of 192)

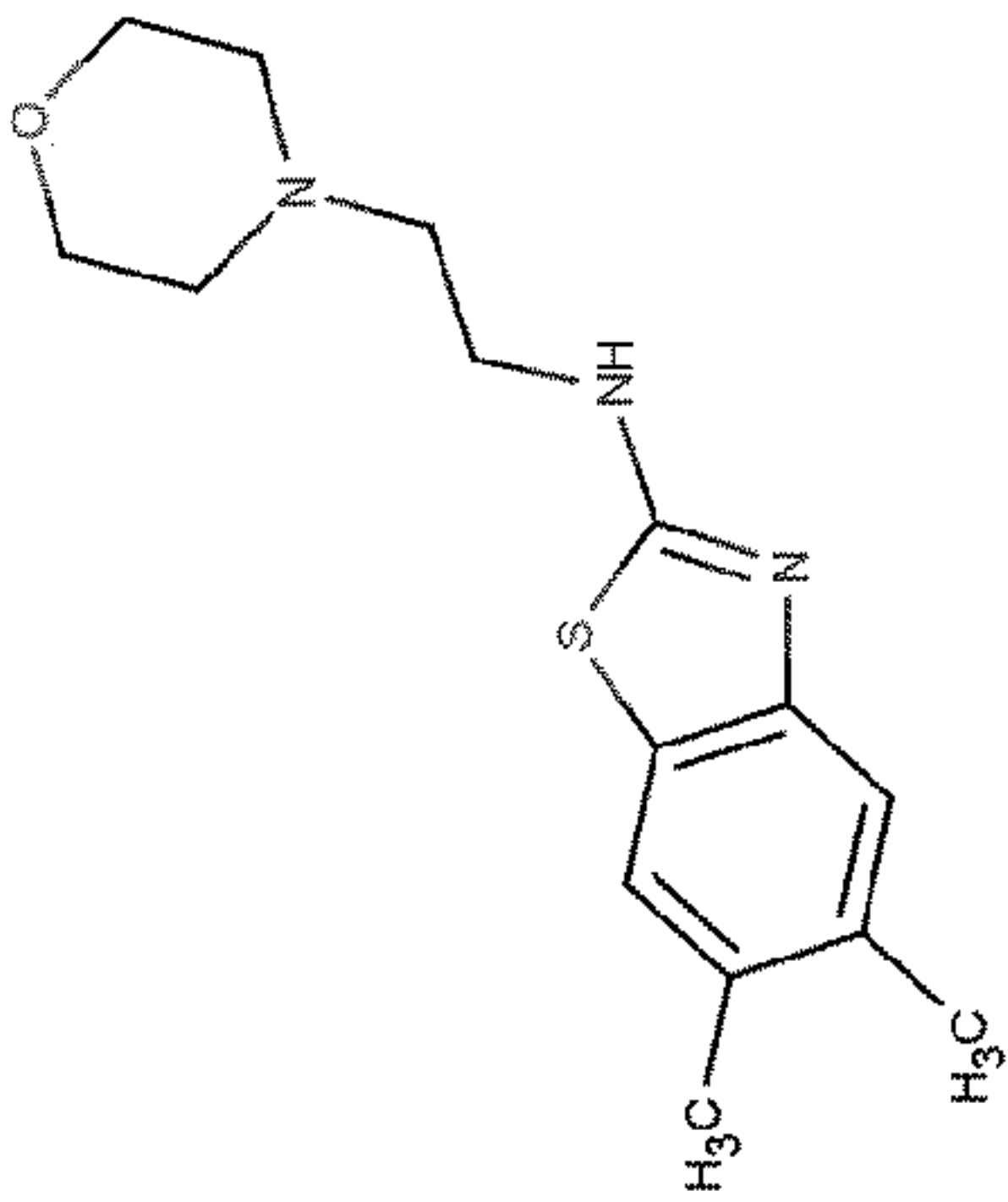
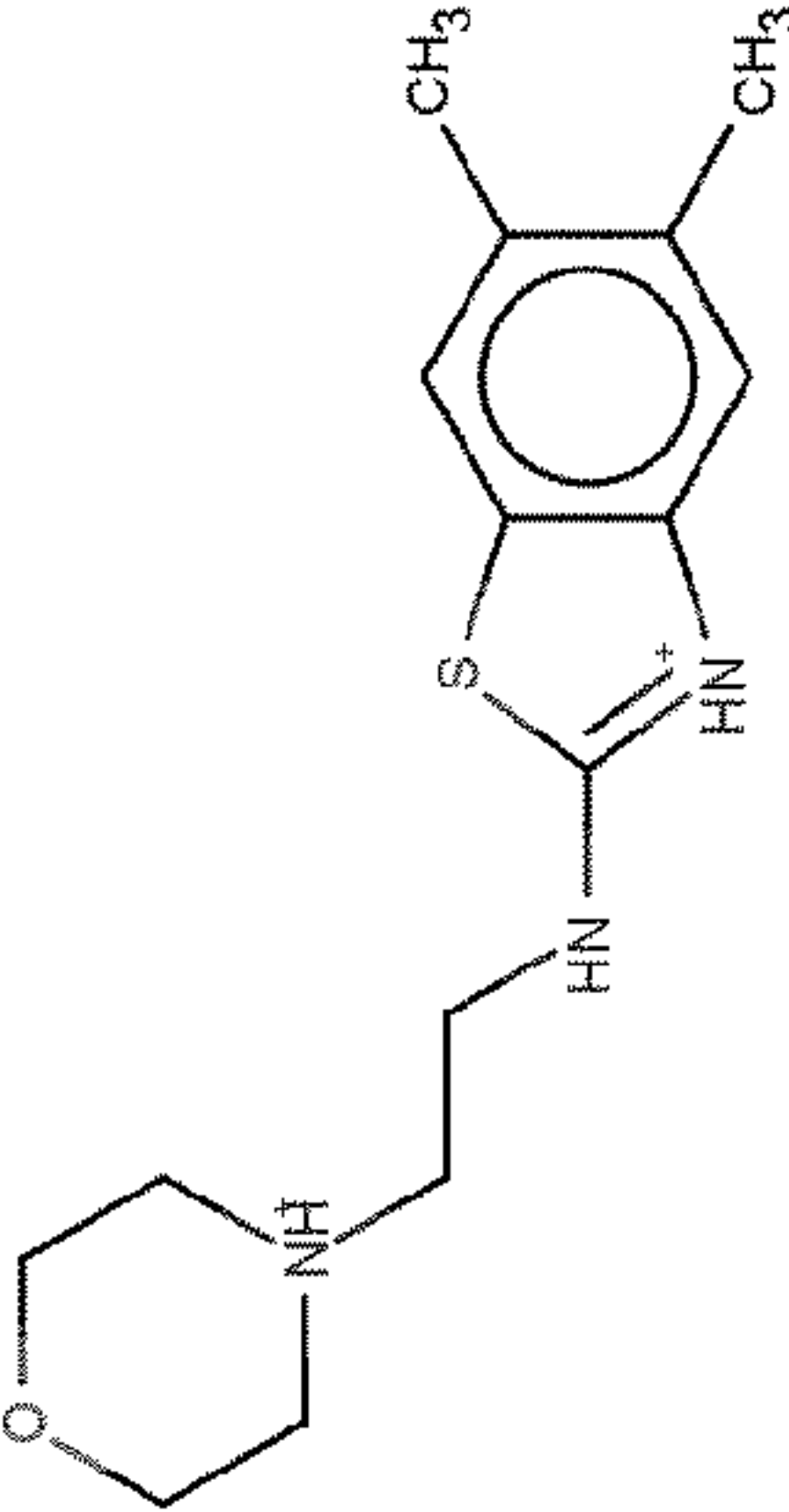
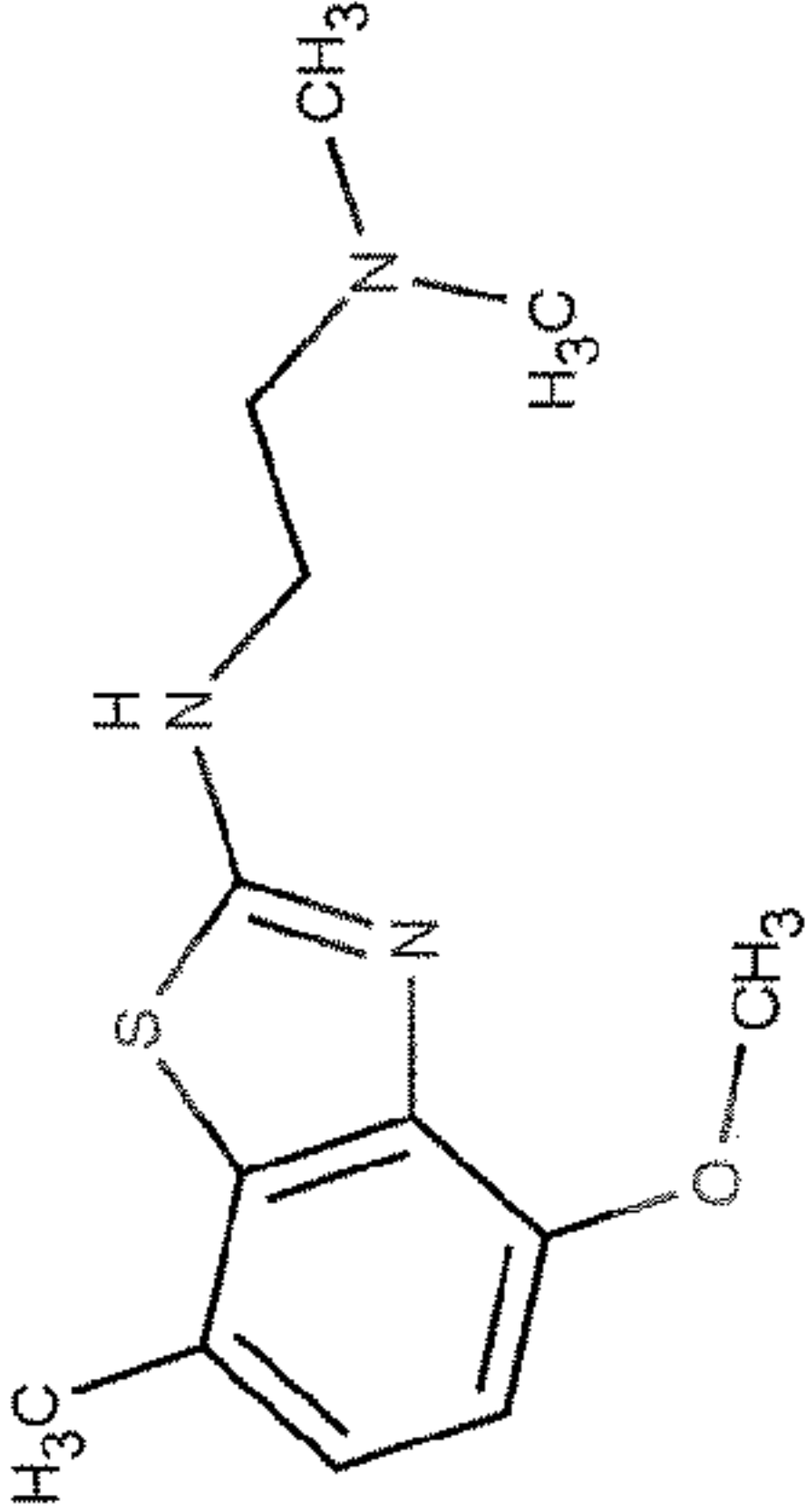
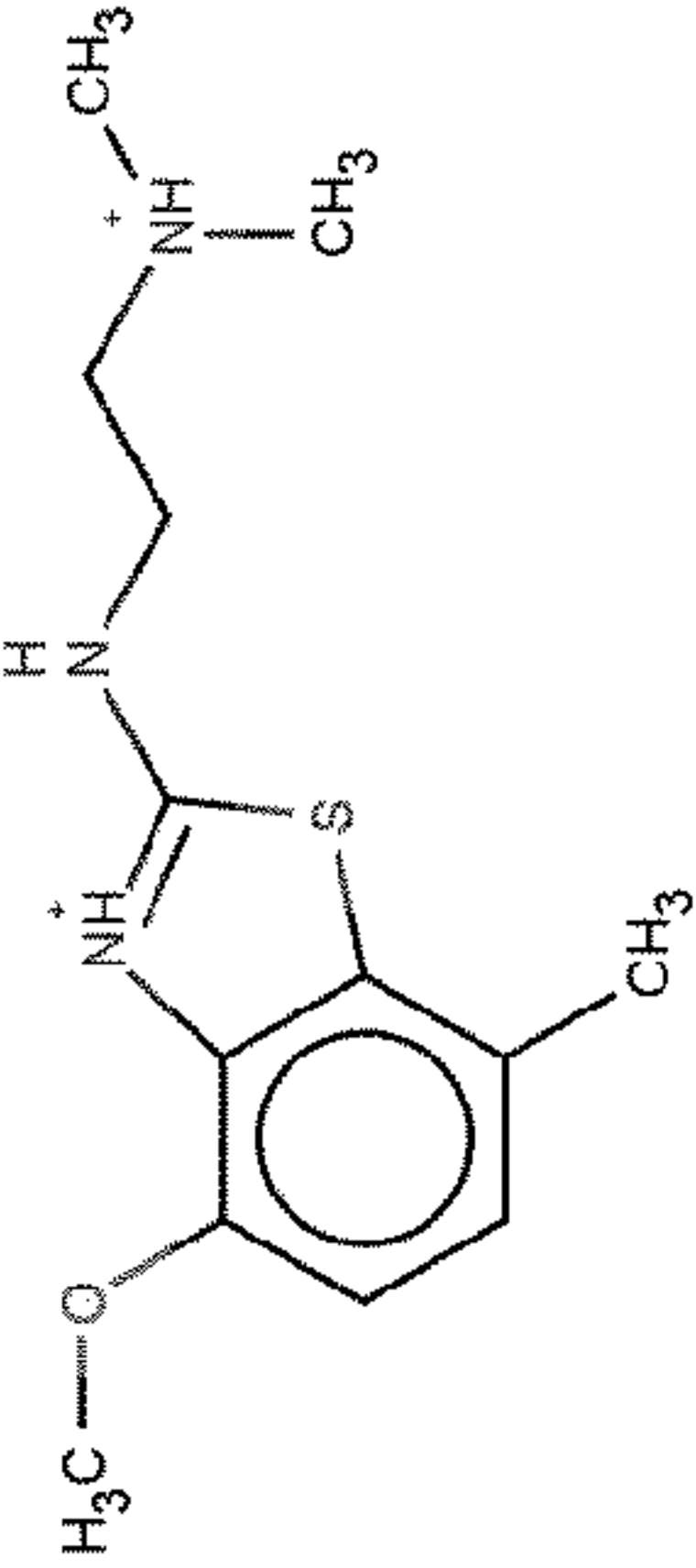
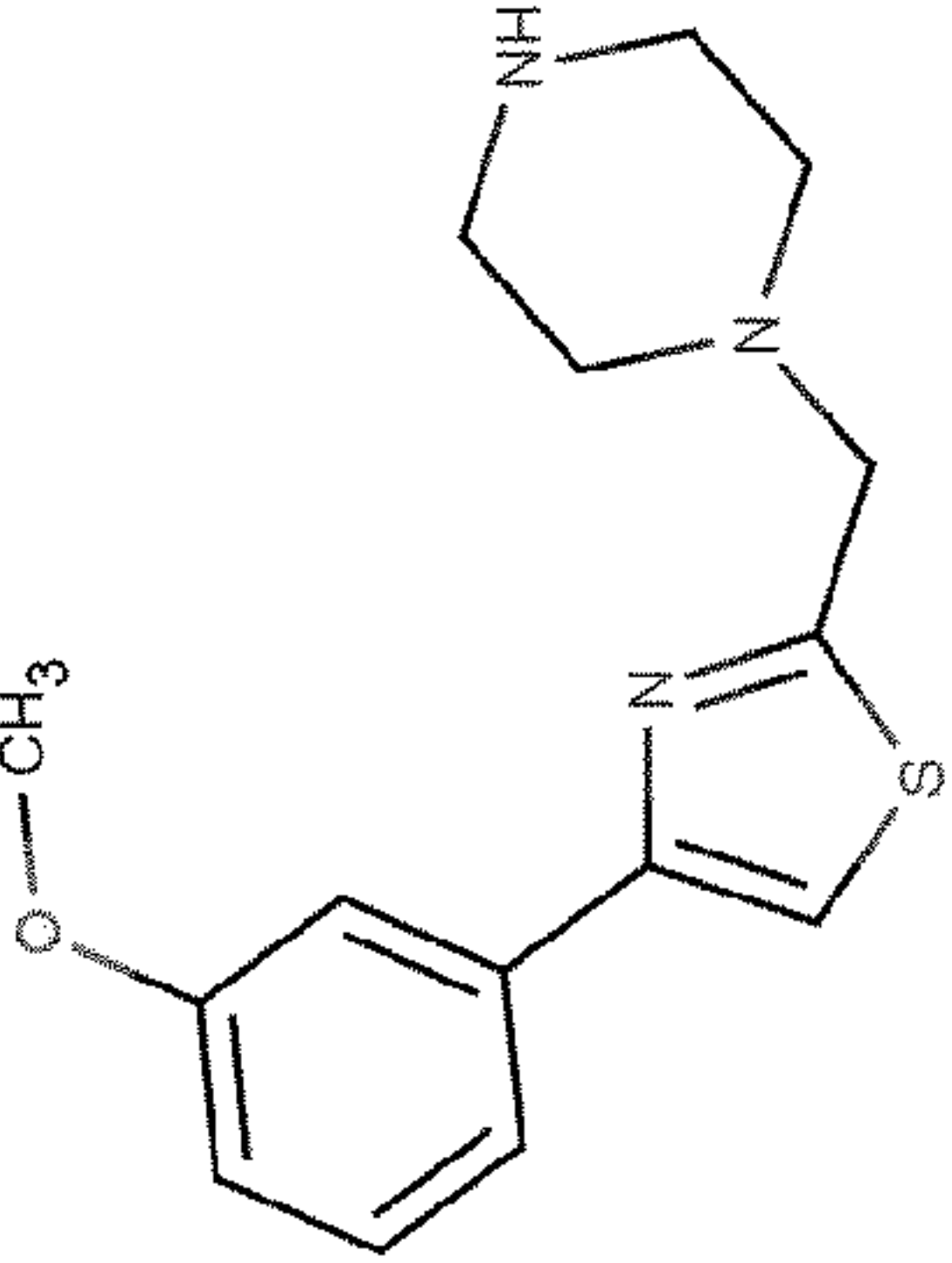
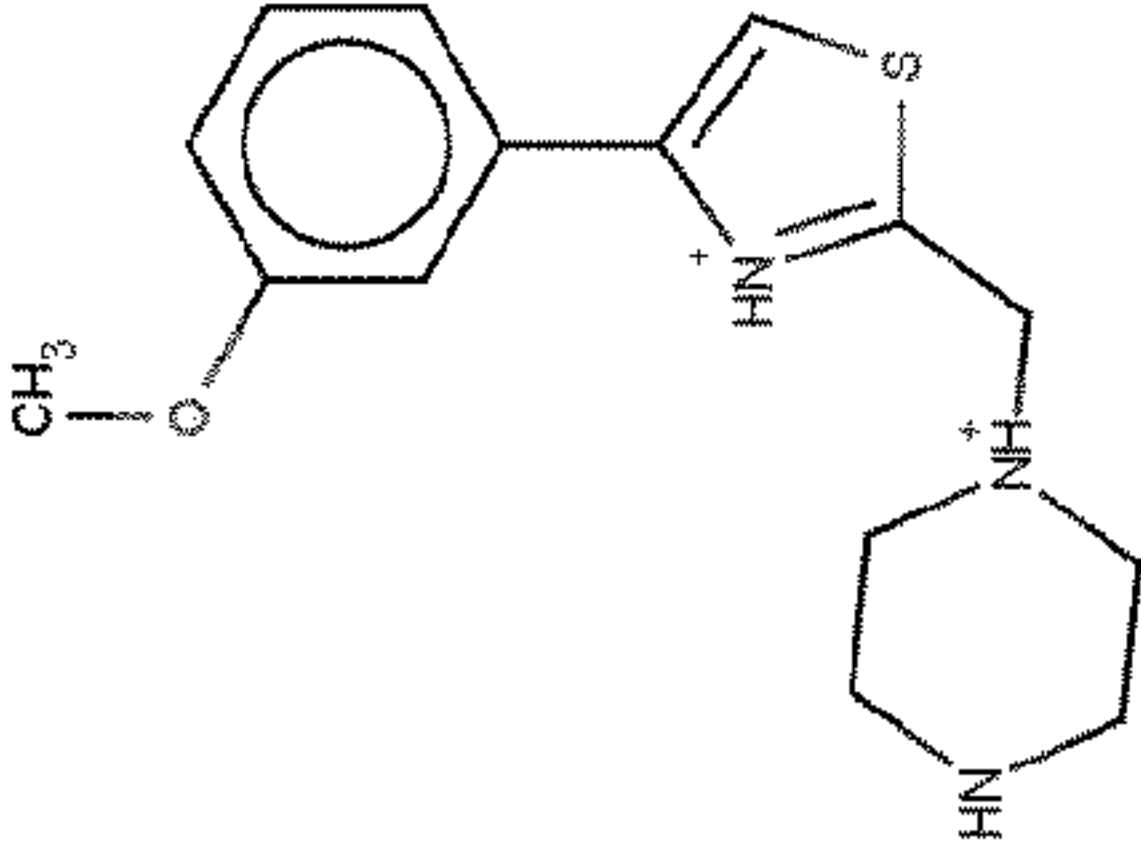
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 81		F1811-0121		F1811-0121_142_1 (1 of 171)
Compound 82		F1863-0017		F1863-0017_2 (1 of 28)
Compound 83		F1894-0028		F1894-0028_3_1 (1 of 6)

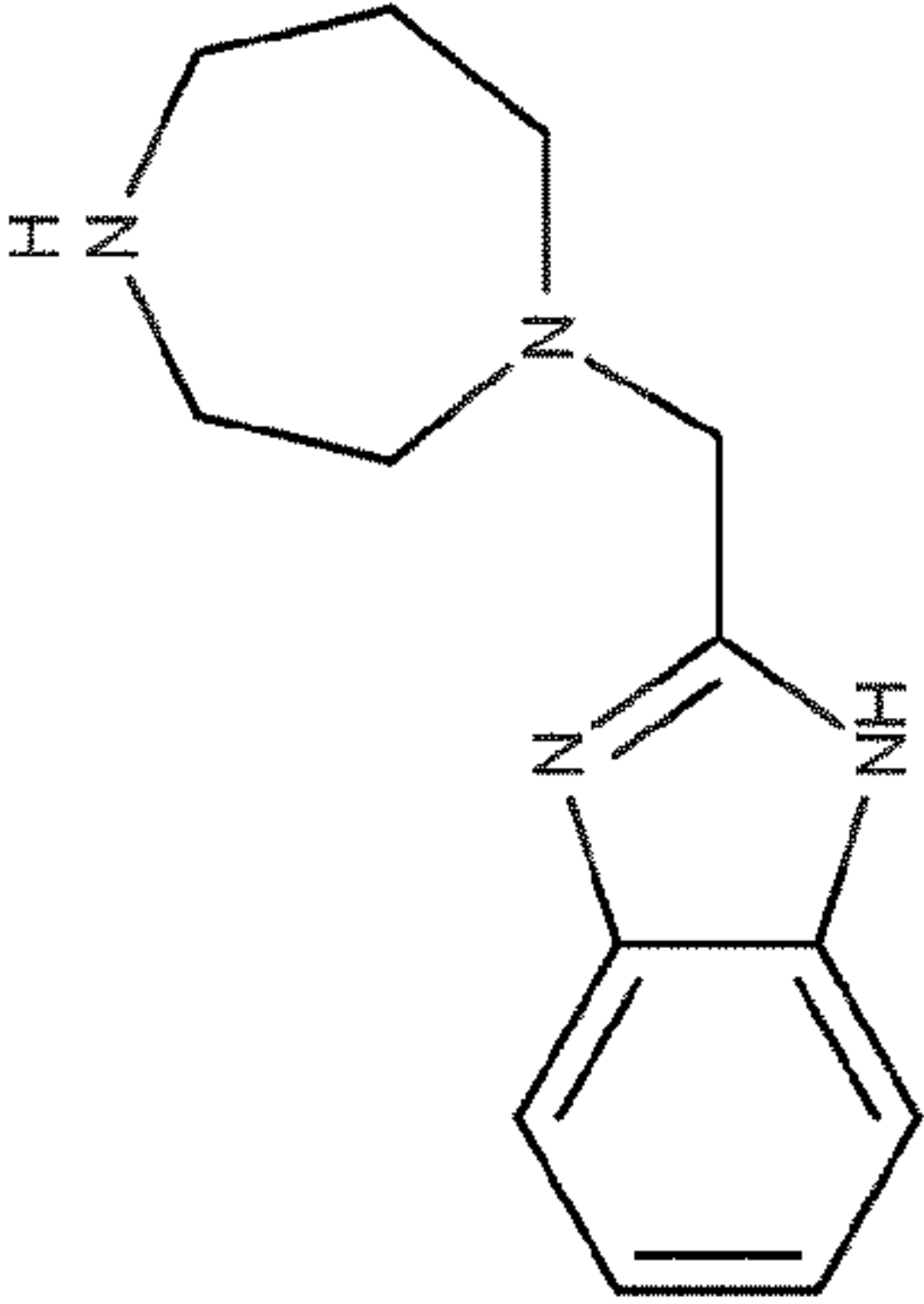
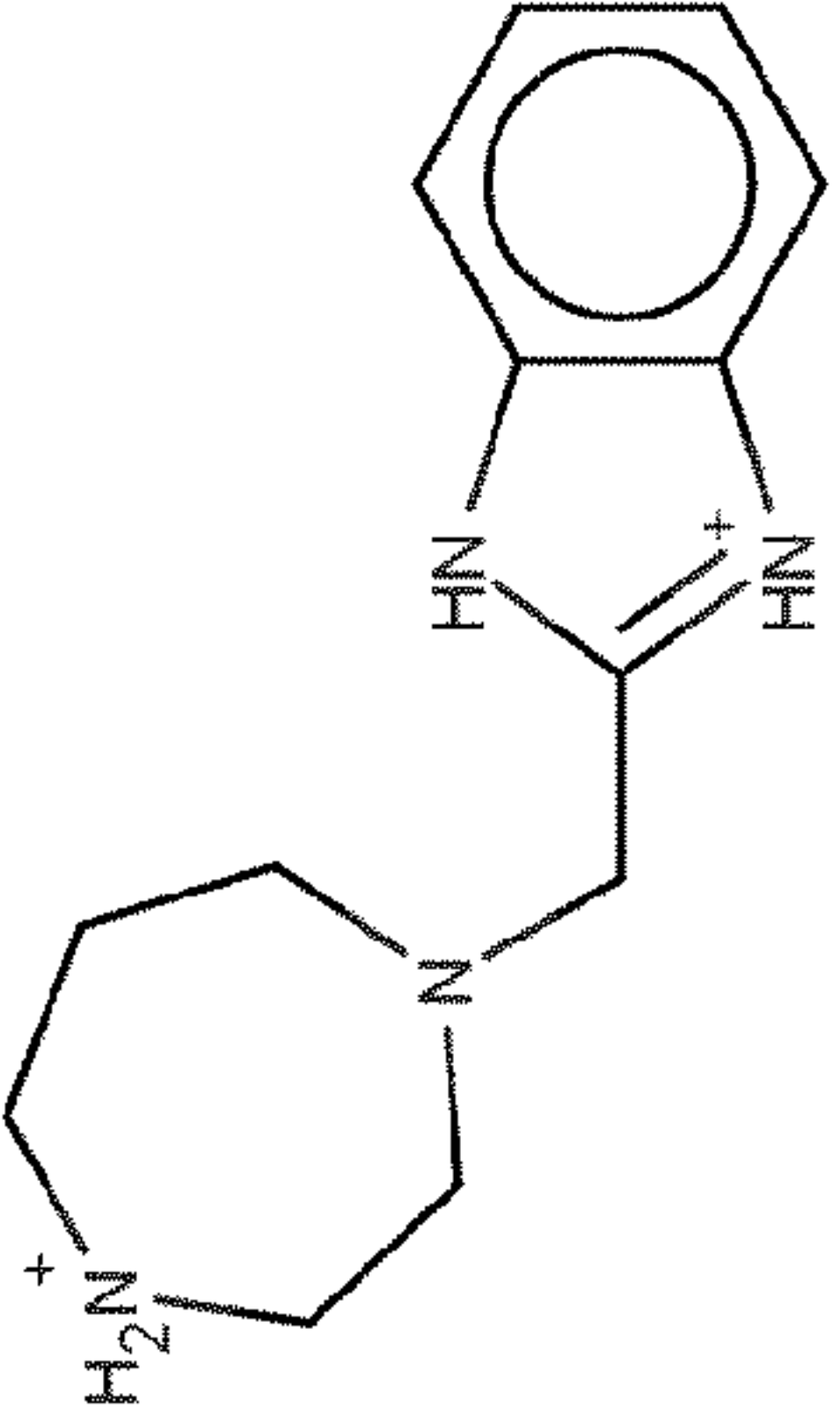
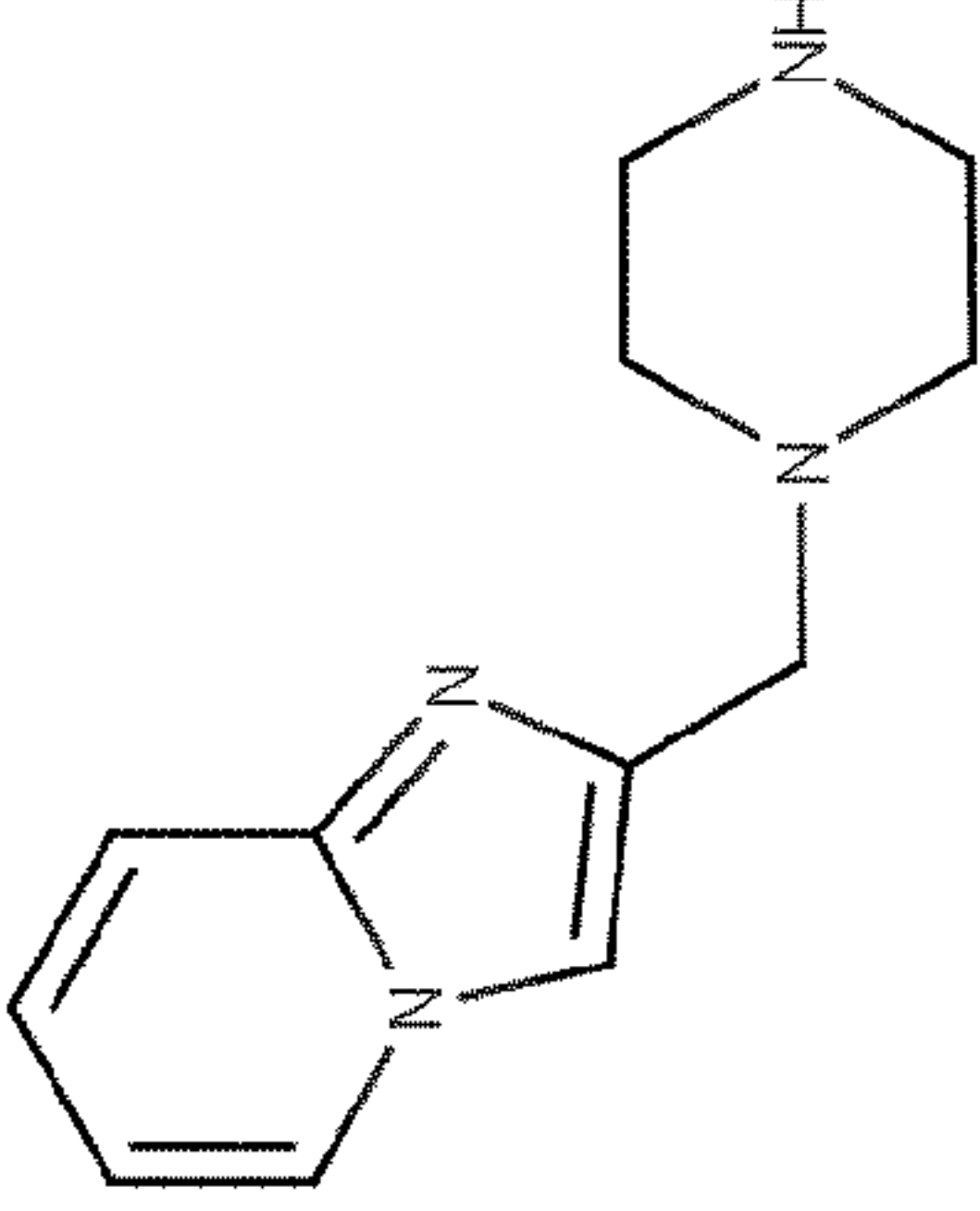
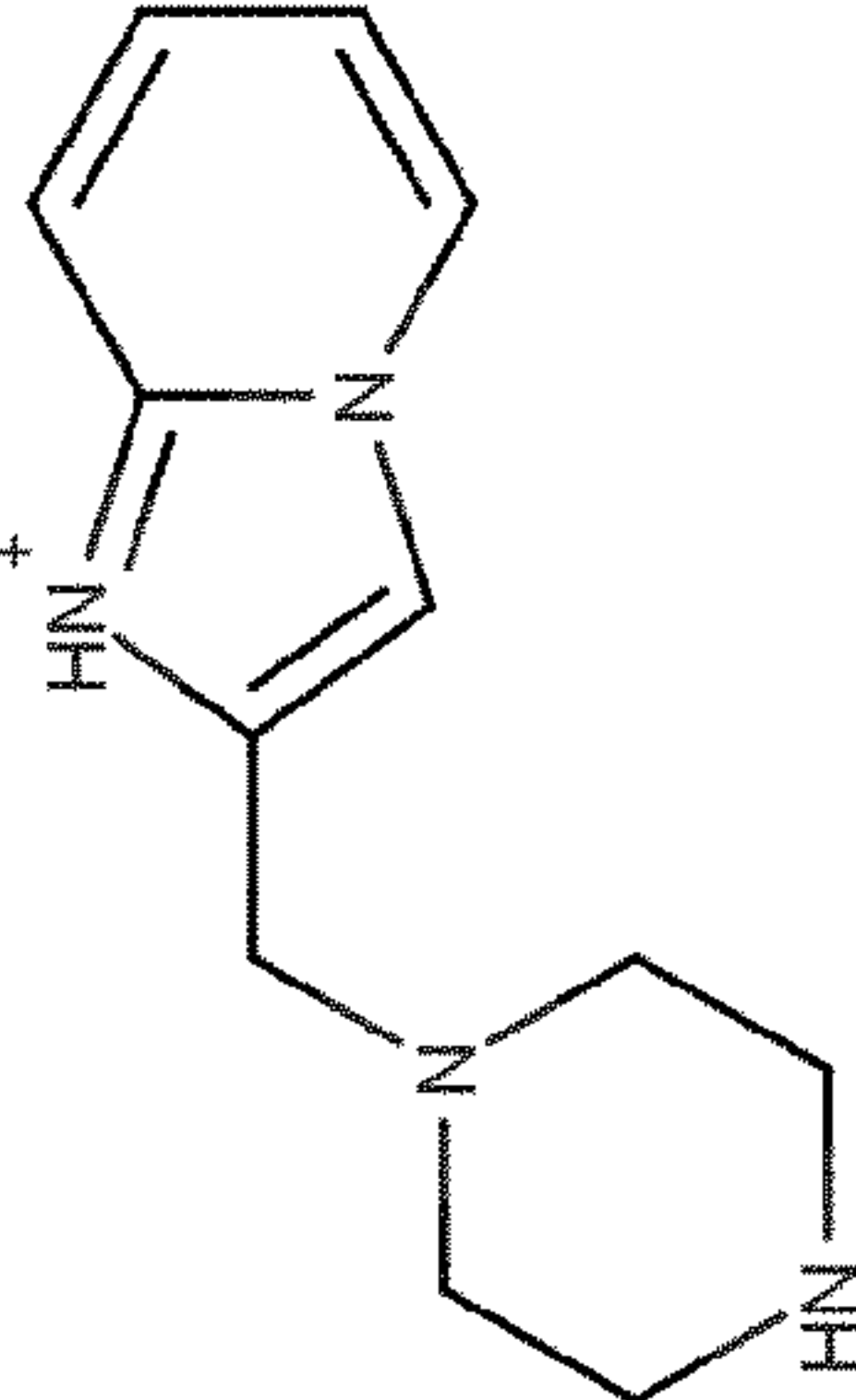
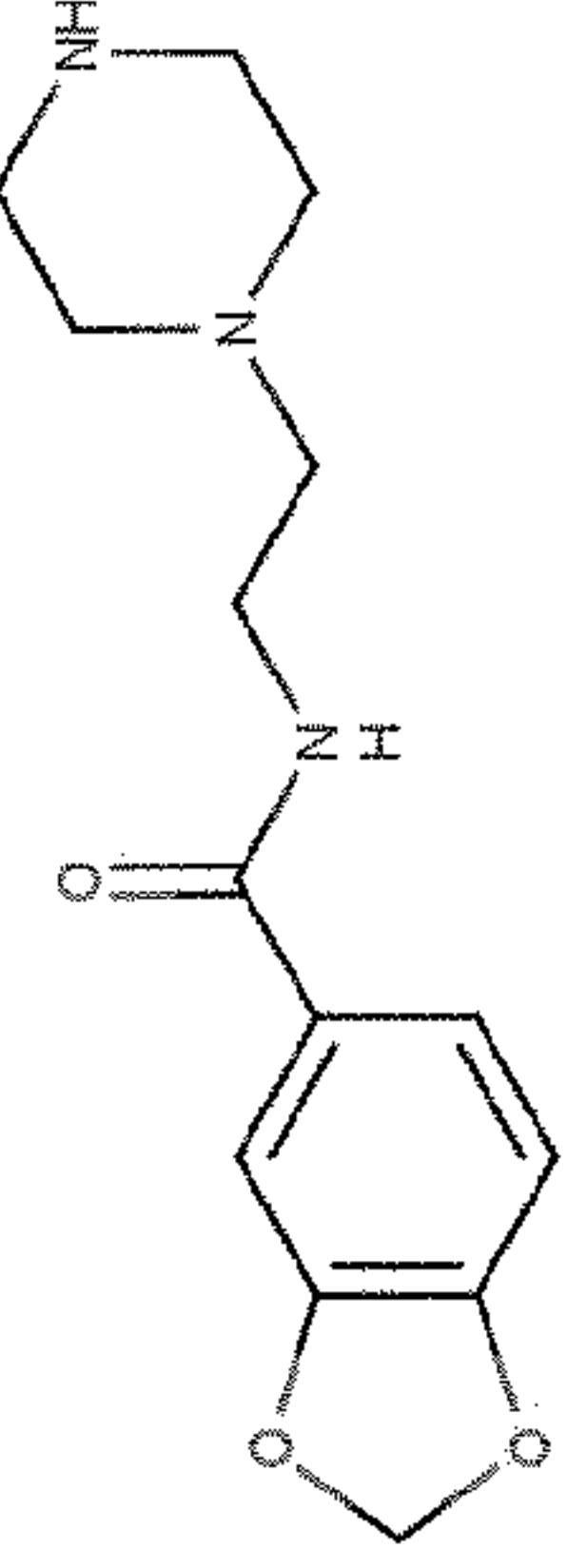
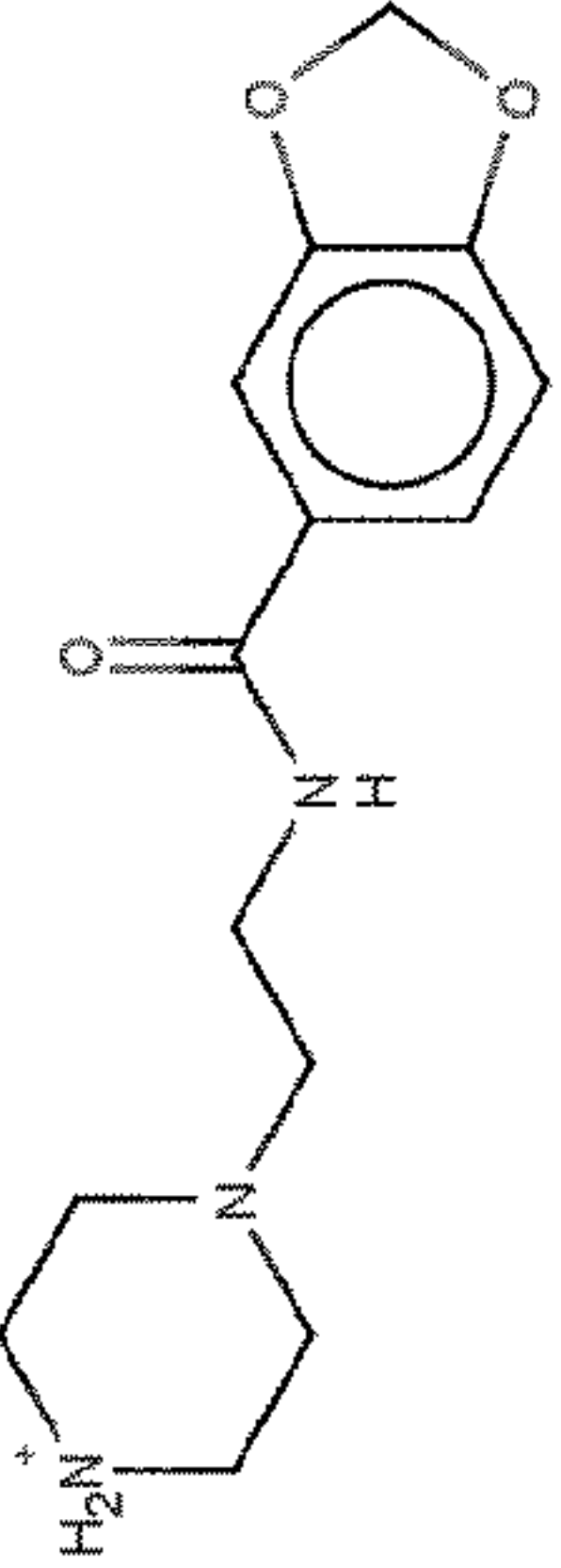
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 84		F1894-0031		F1894-0031_3_1 (1 of 3)
Compound 85		F1894-0157		F1894-0157_3_1 (1 of 5)
Compound 86		F1967-1078		F1967-1078_1 (1 of 4)

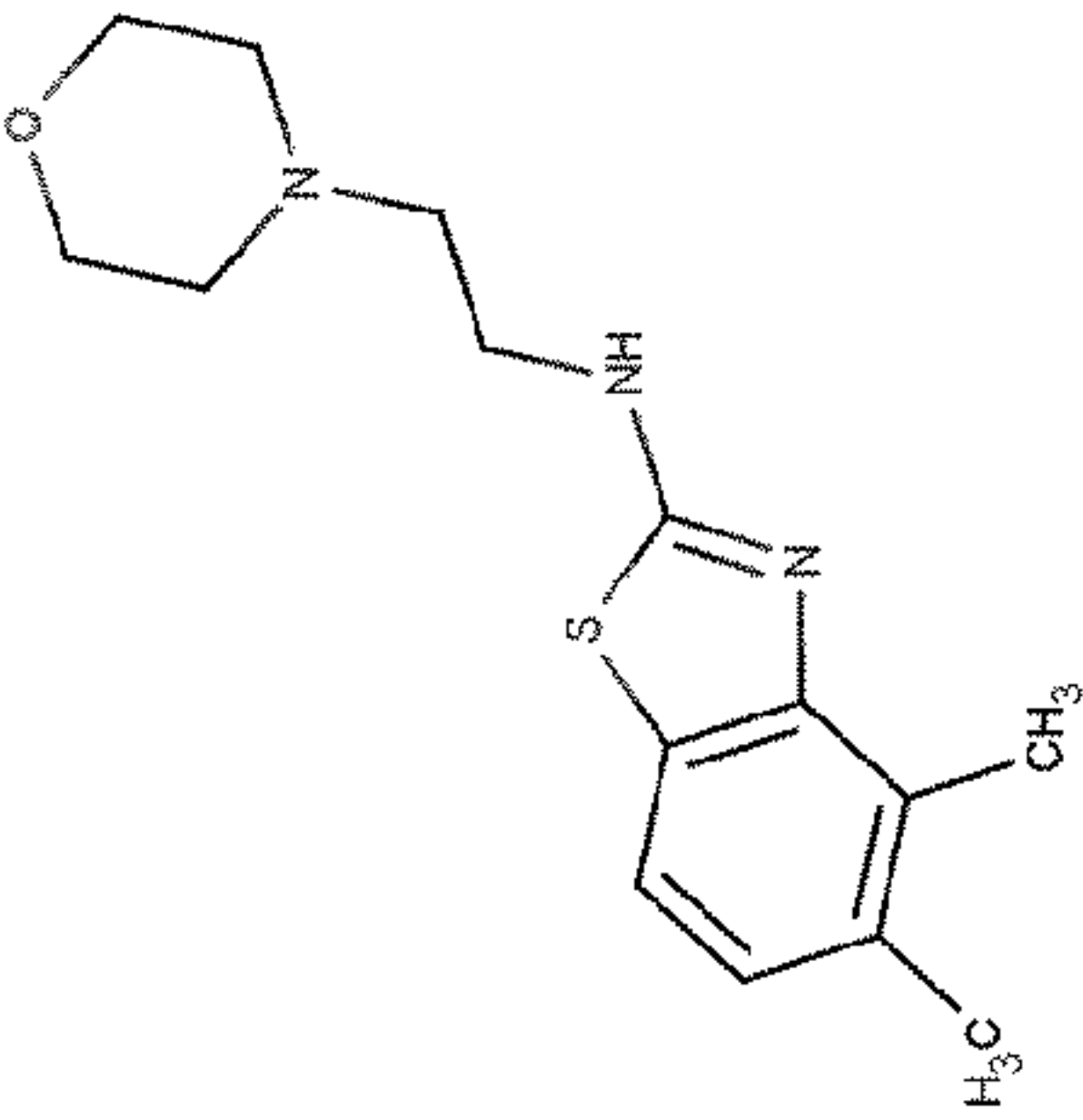
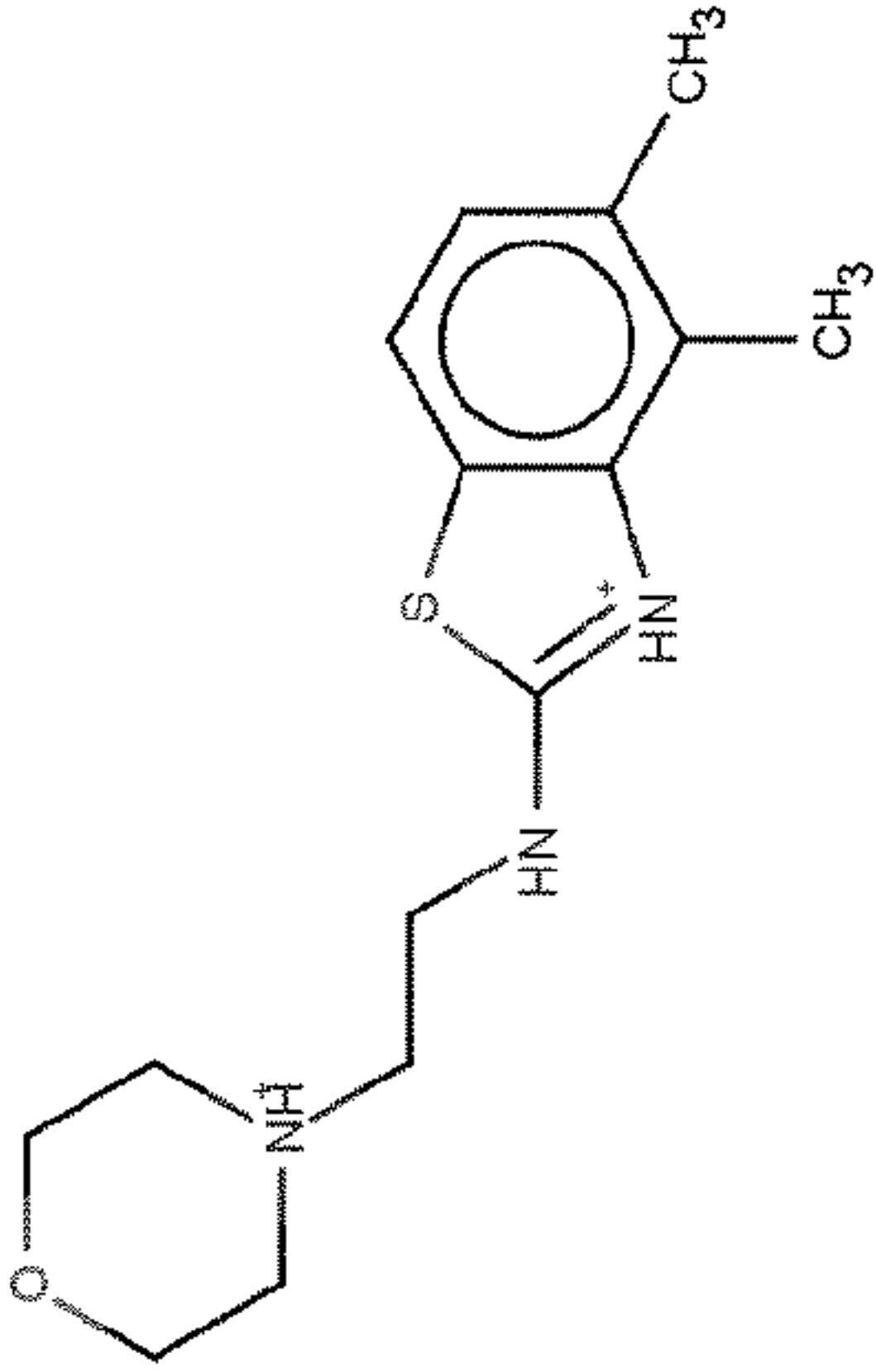
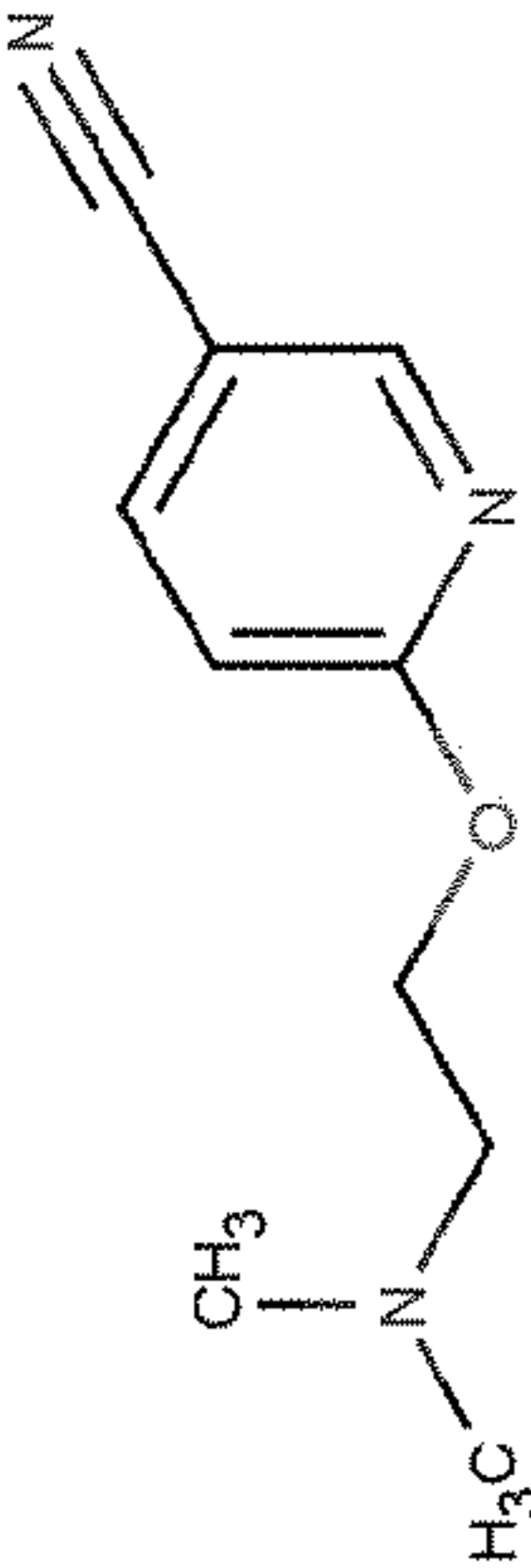
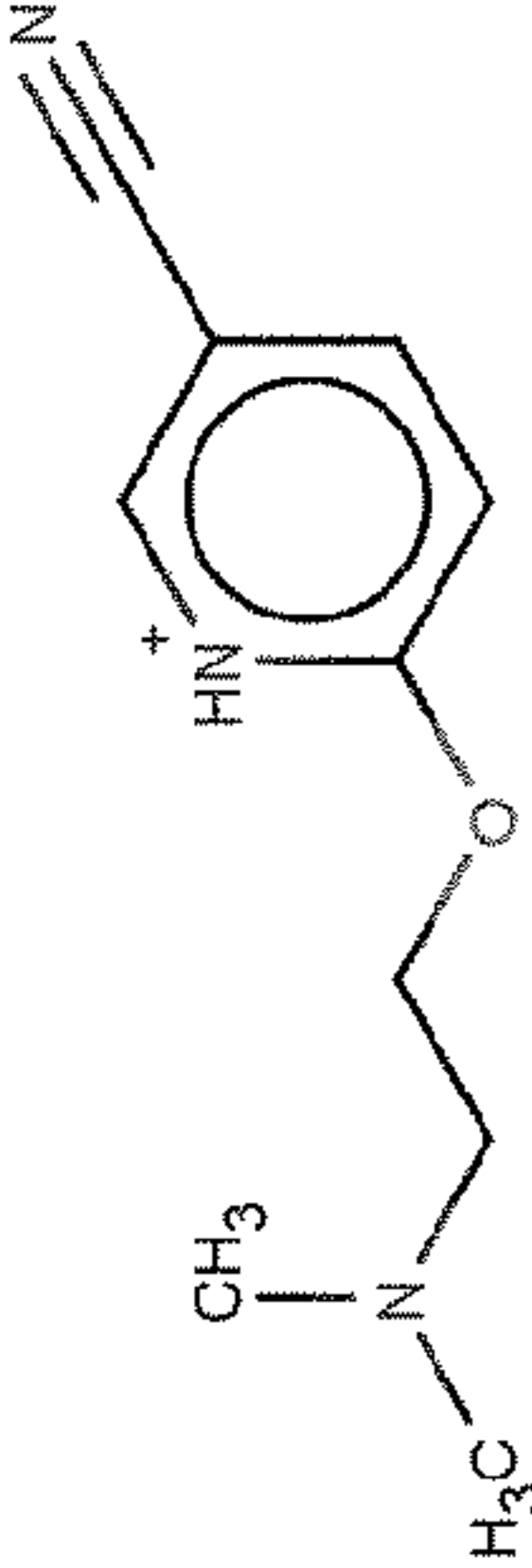
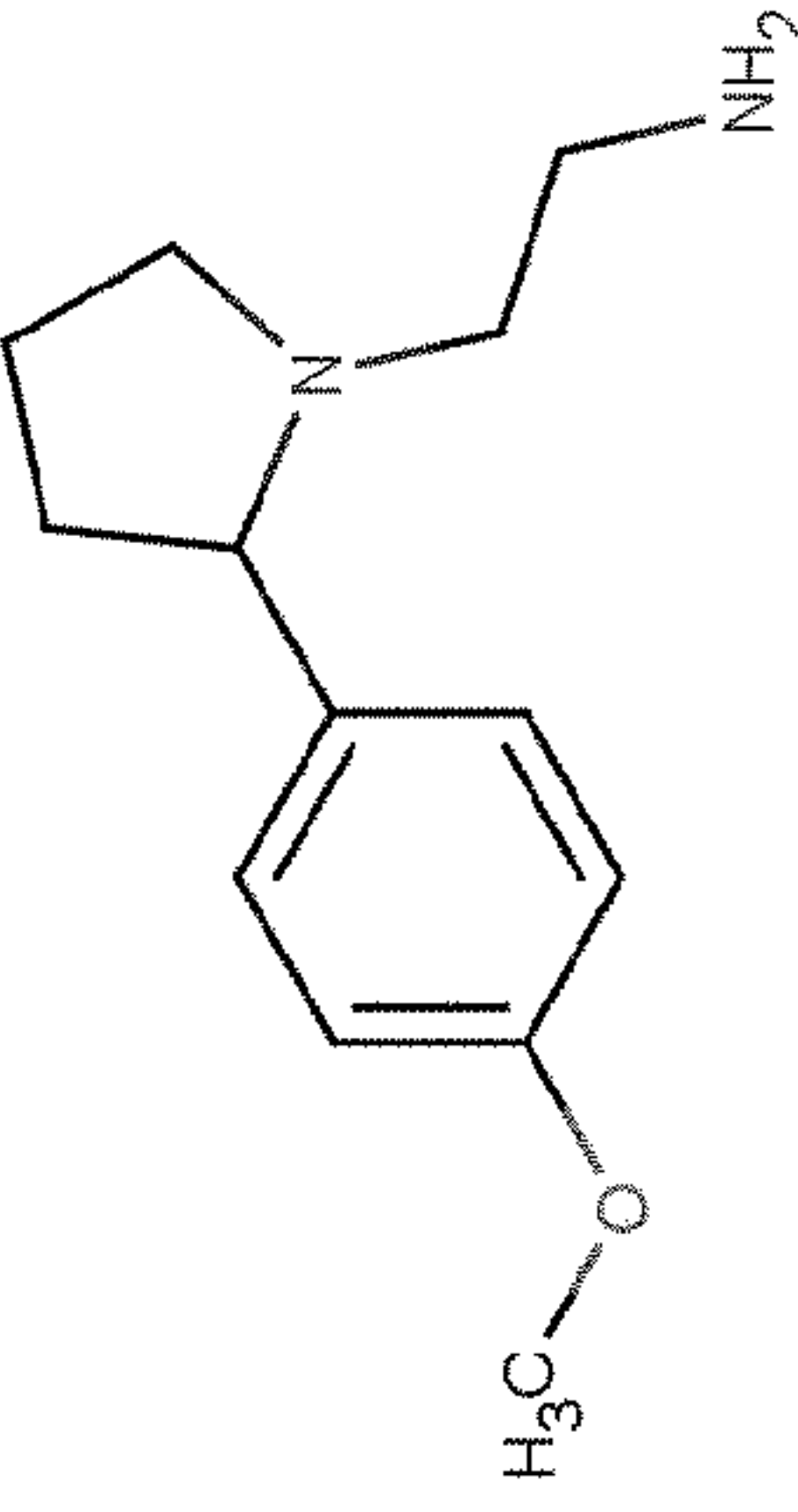
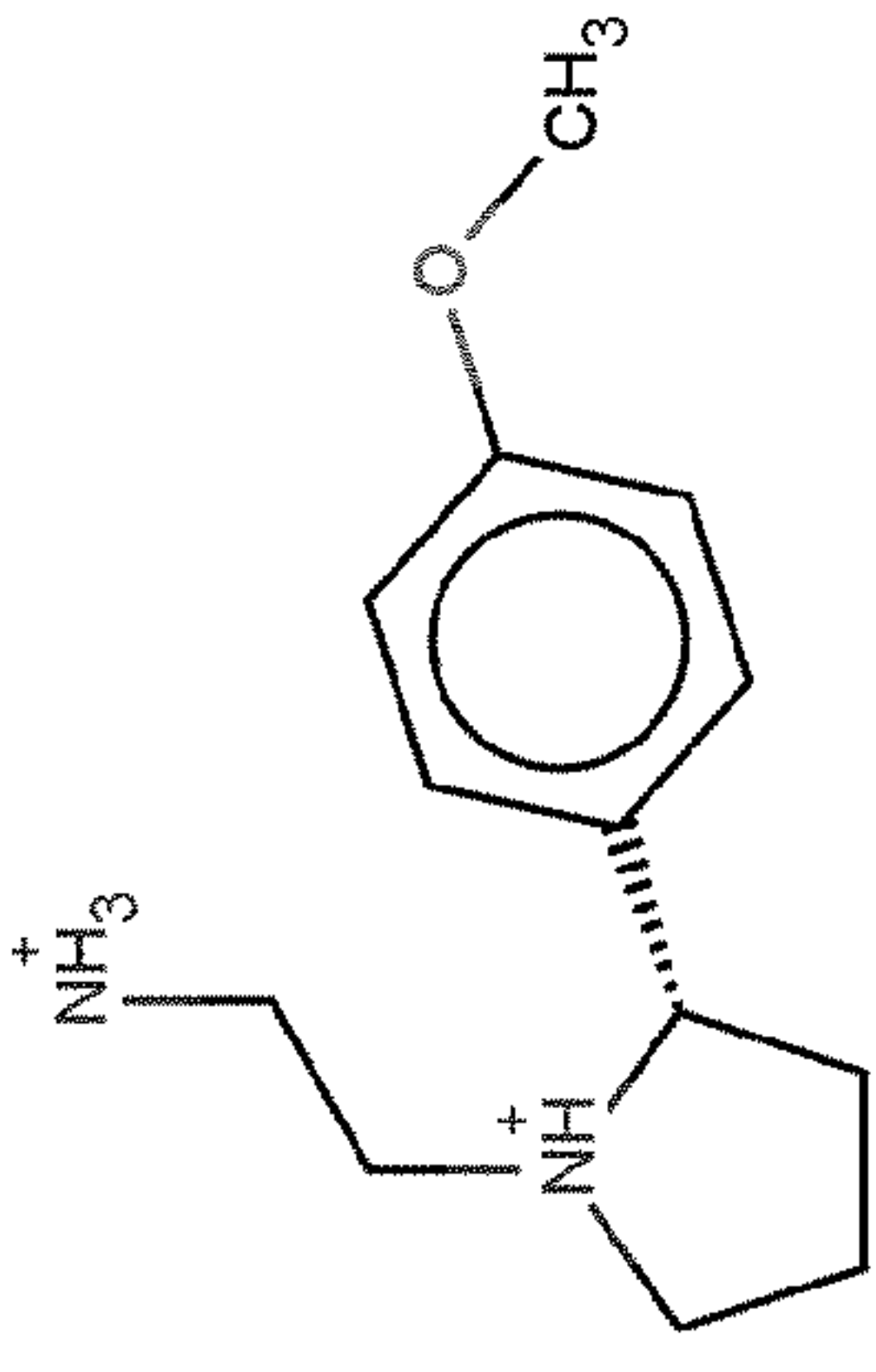
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 87		F1967-1118		F1967-1118_22_1 (1 of 41)
Compound 88		F1967-1334		F1967-1334_1_1 (1 of 2)
Compound 89		F1984-0137		F1984-0137_3_1 (1 of 4)

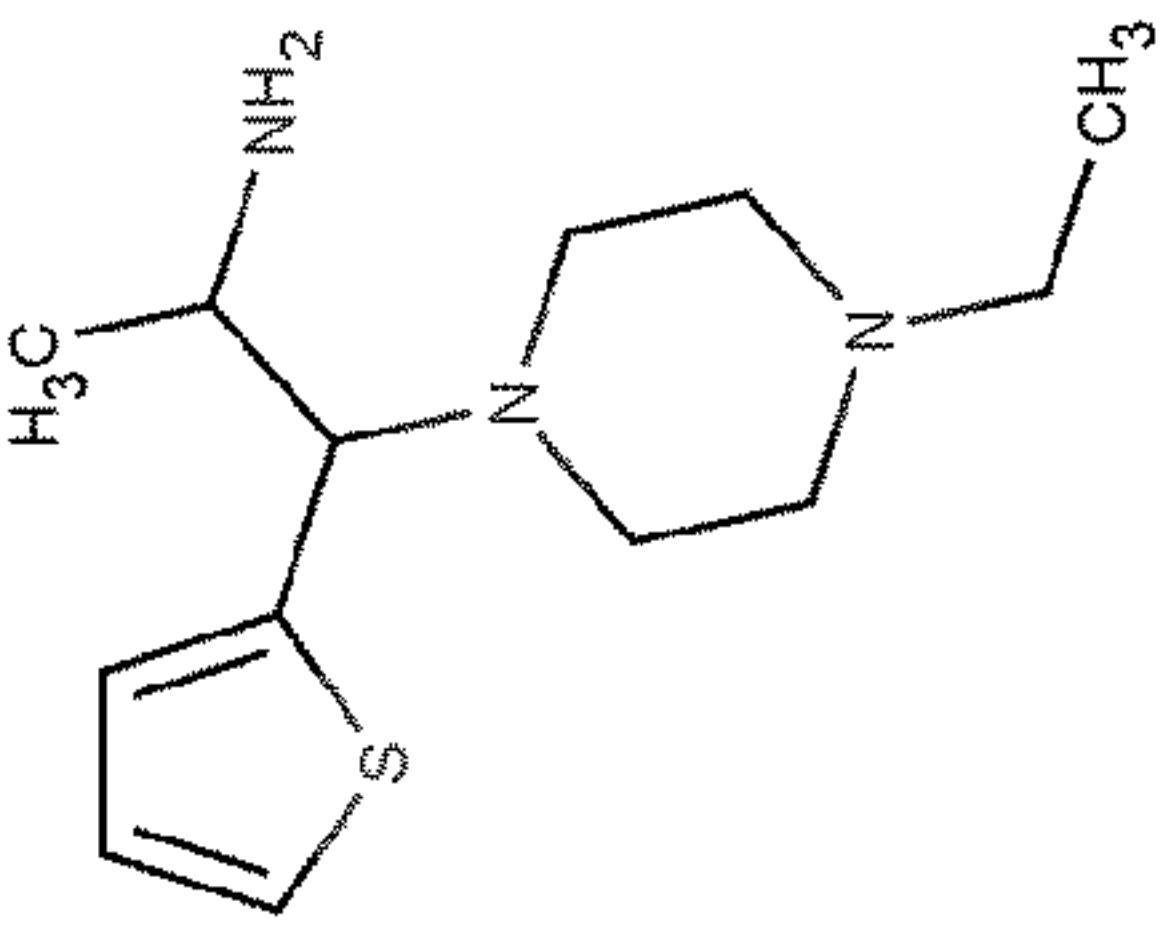
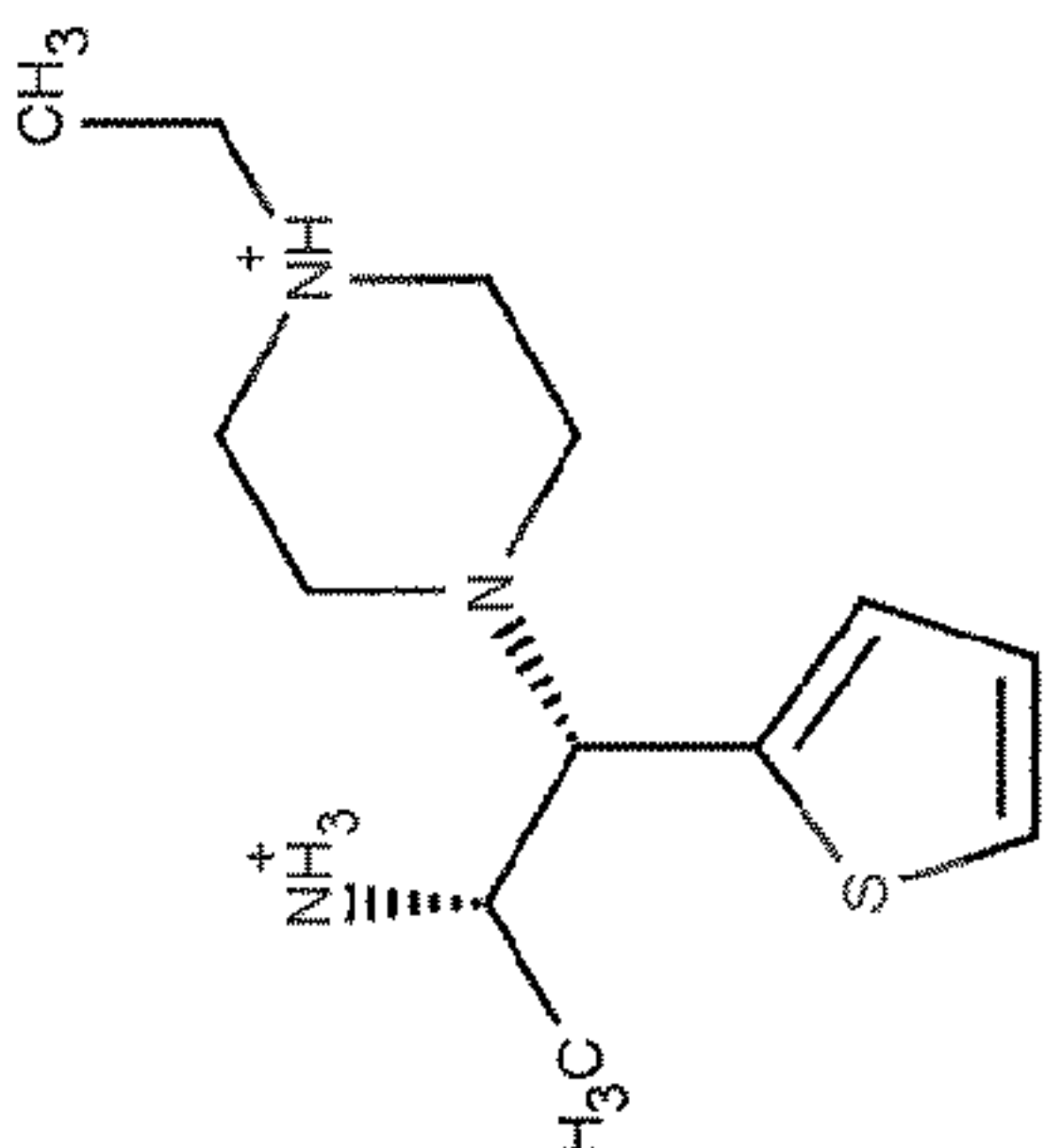
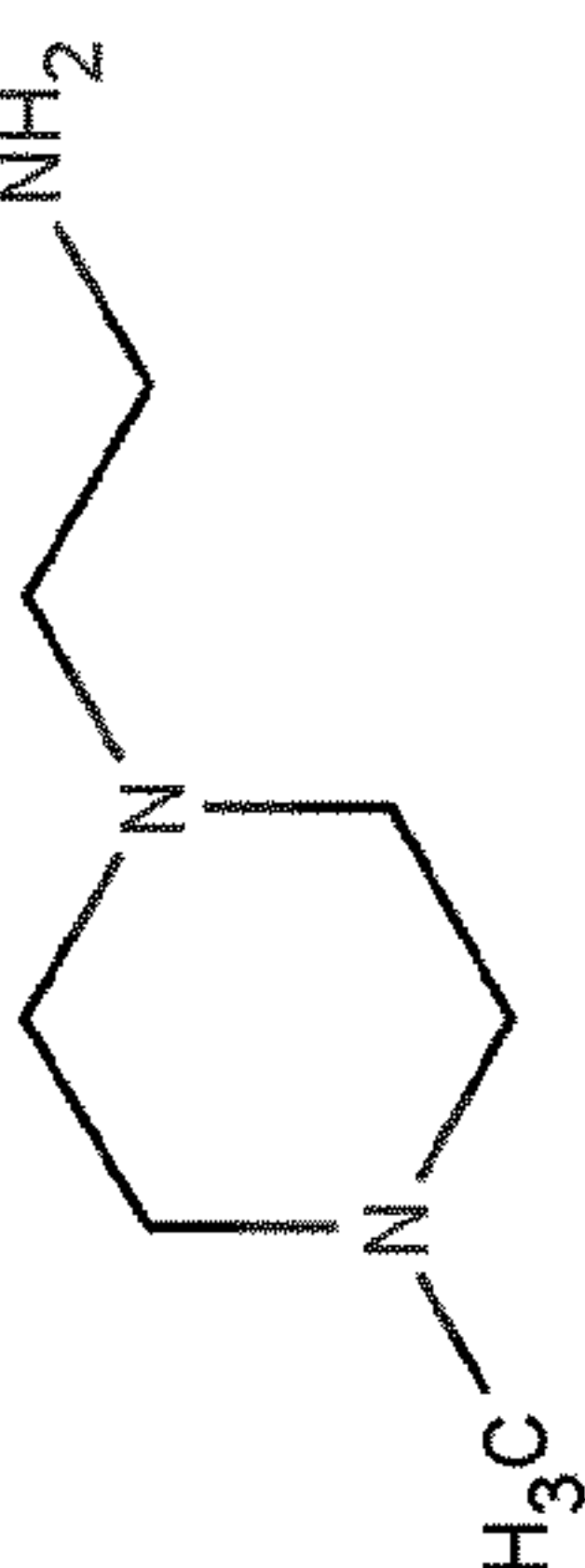
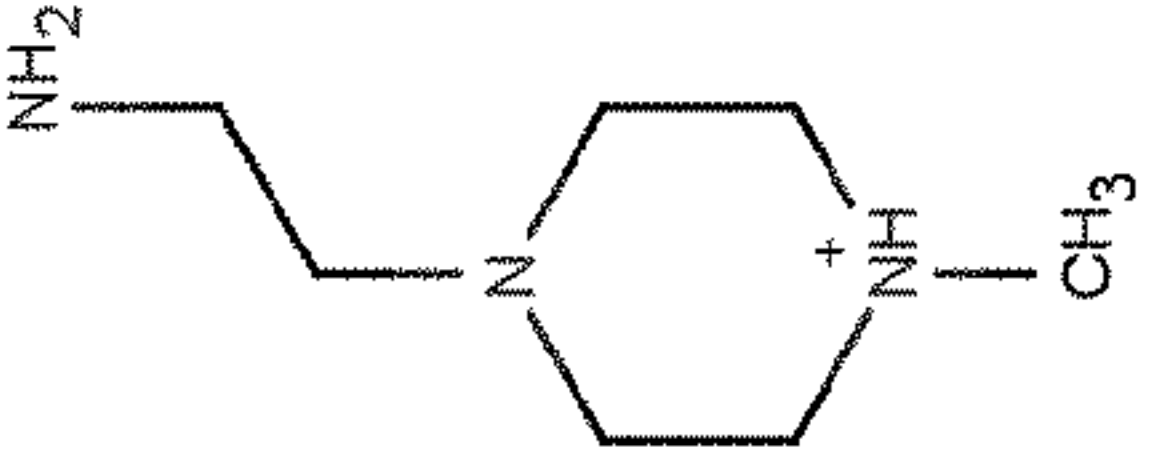
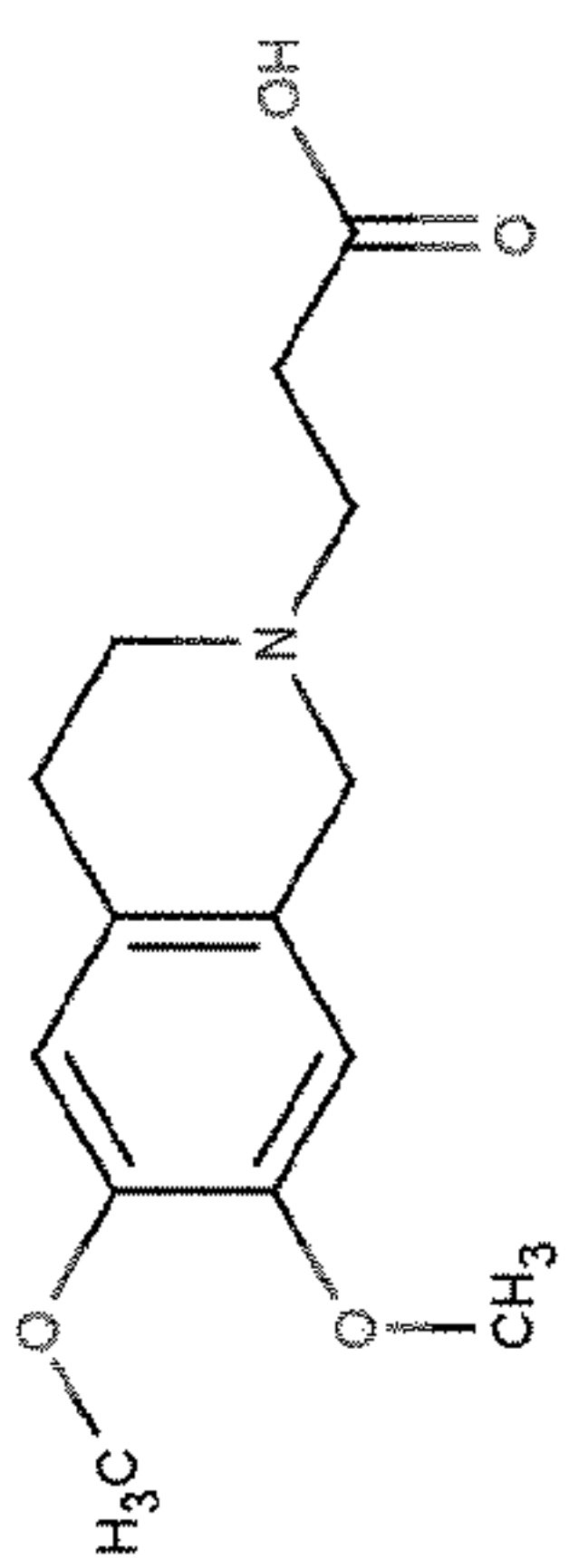
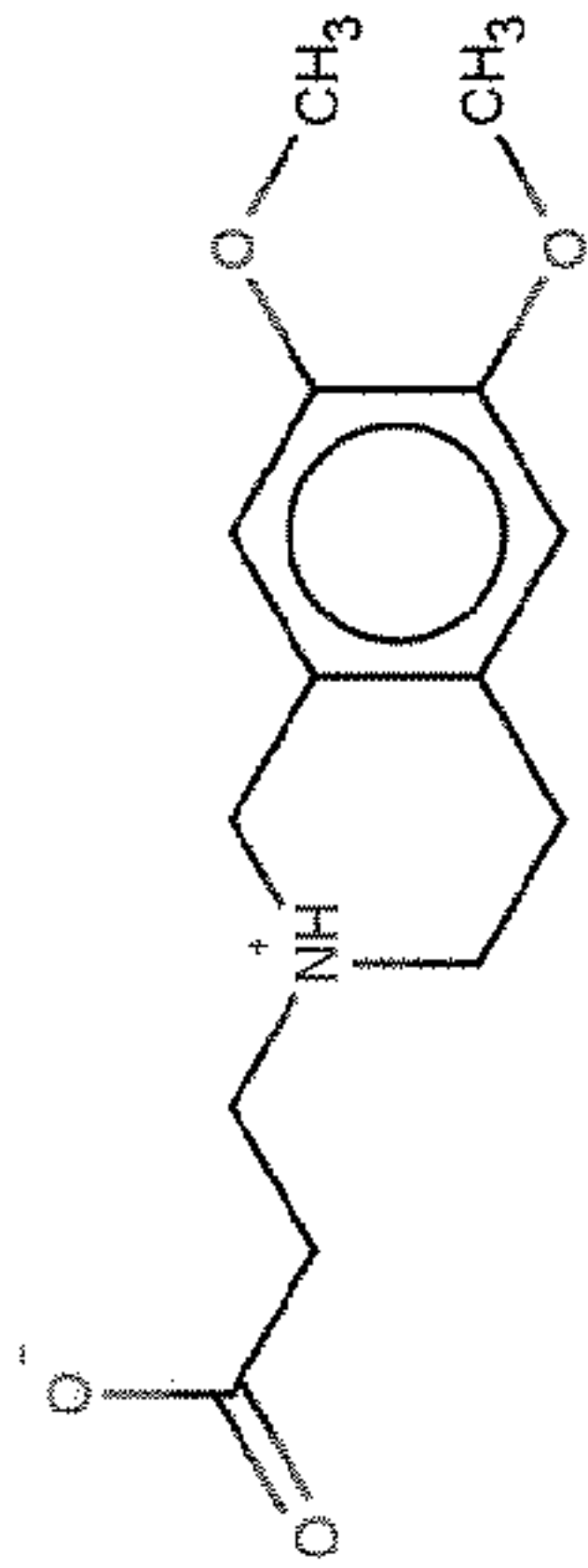
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 90		F1986-0021		F1986-0021_1 (1 of 4)
Compound 91		F1986-0024		F1986-0024_1_1 (1 of 4)
Compound 92		F2011-0385		F2011-0385_1 (1 of 2)

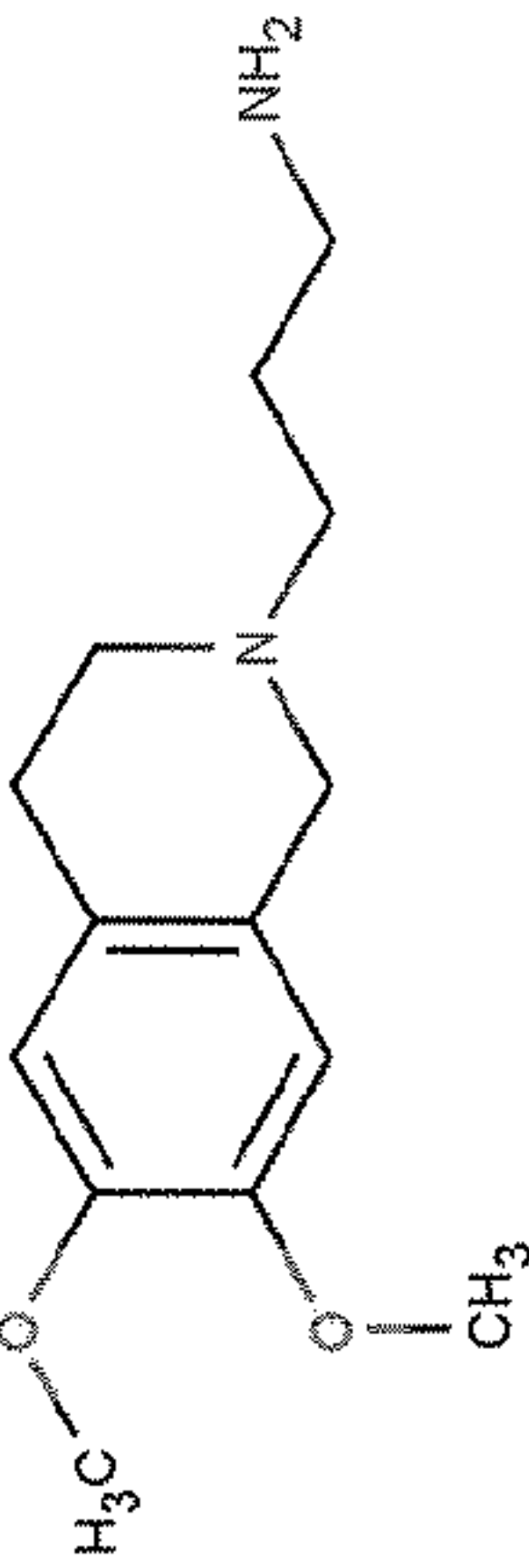
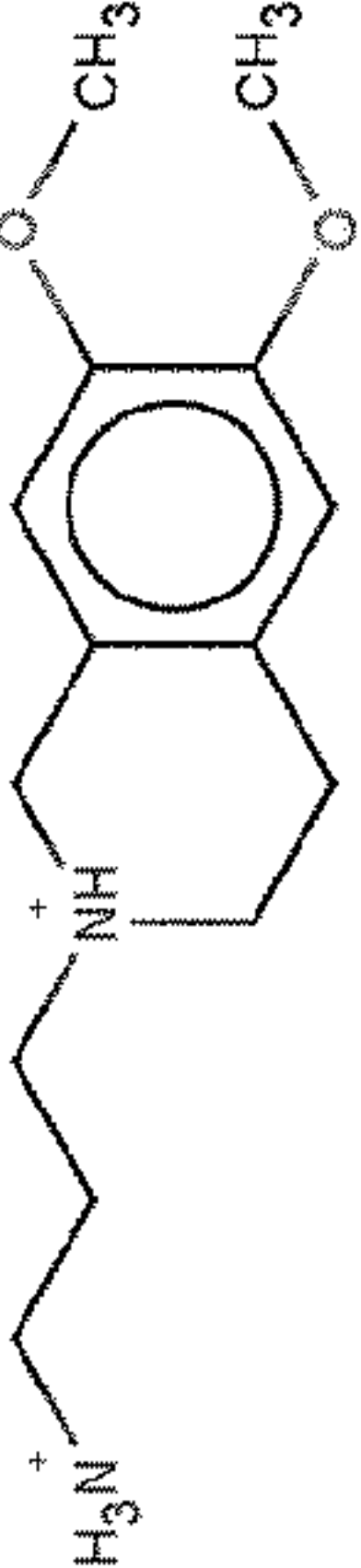
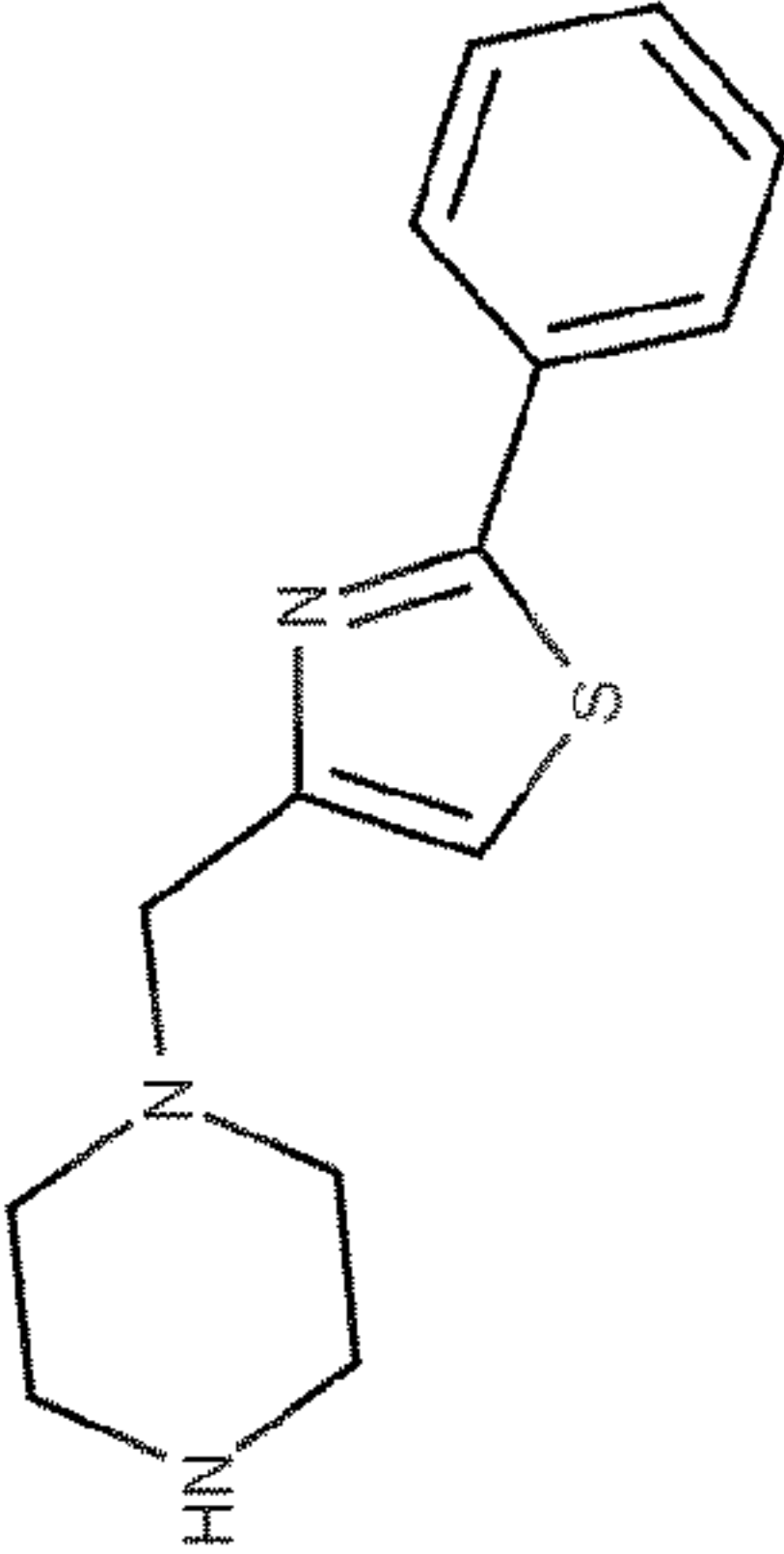
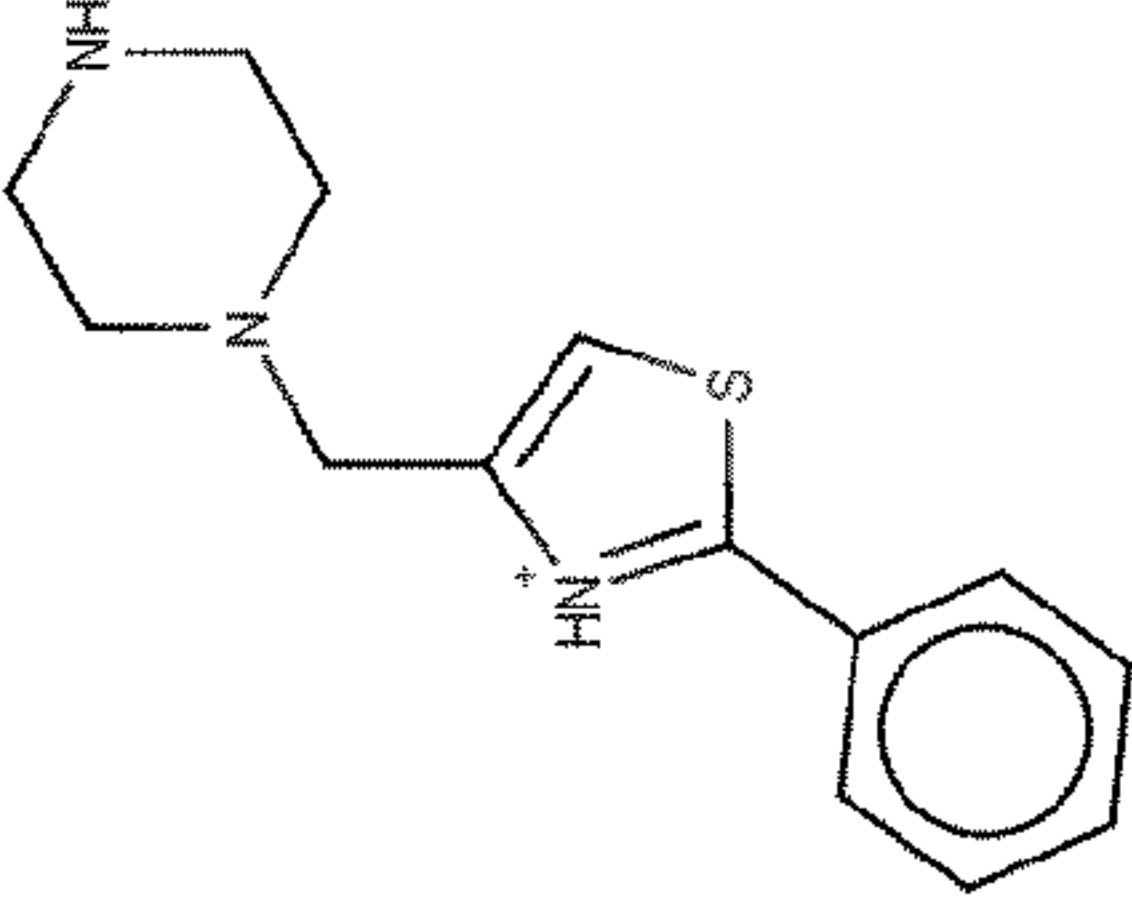
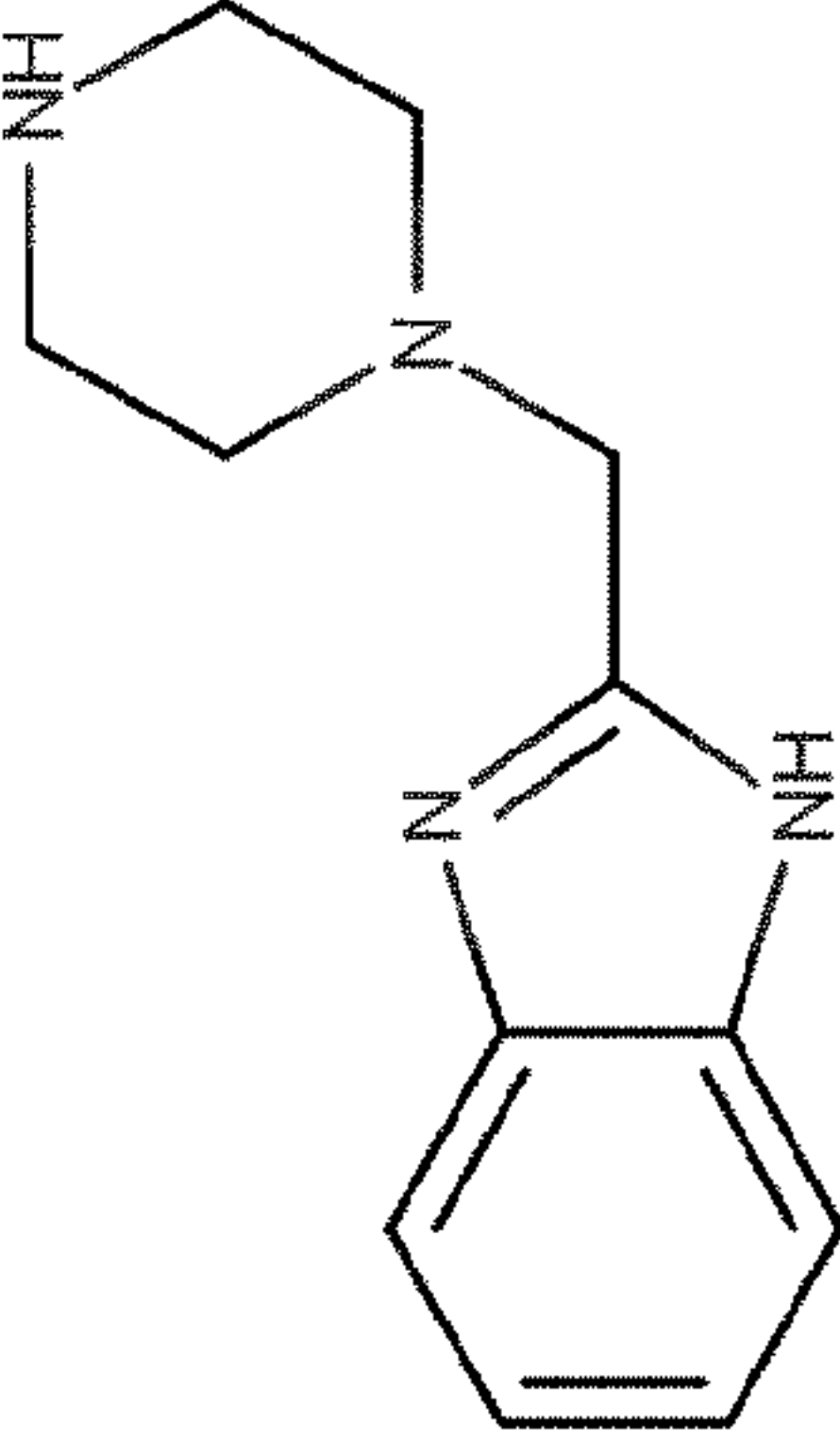
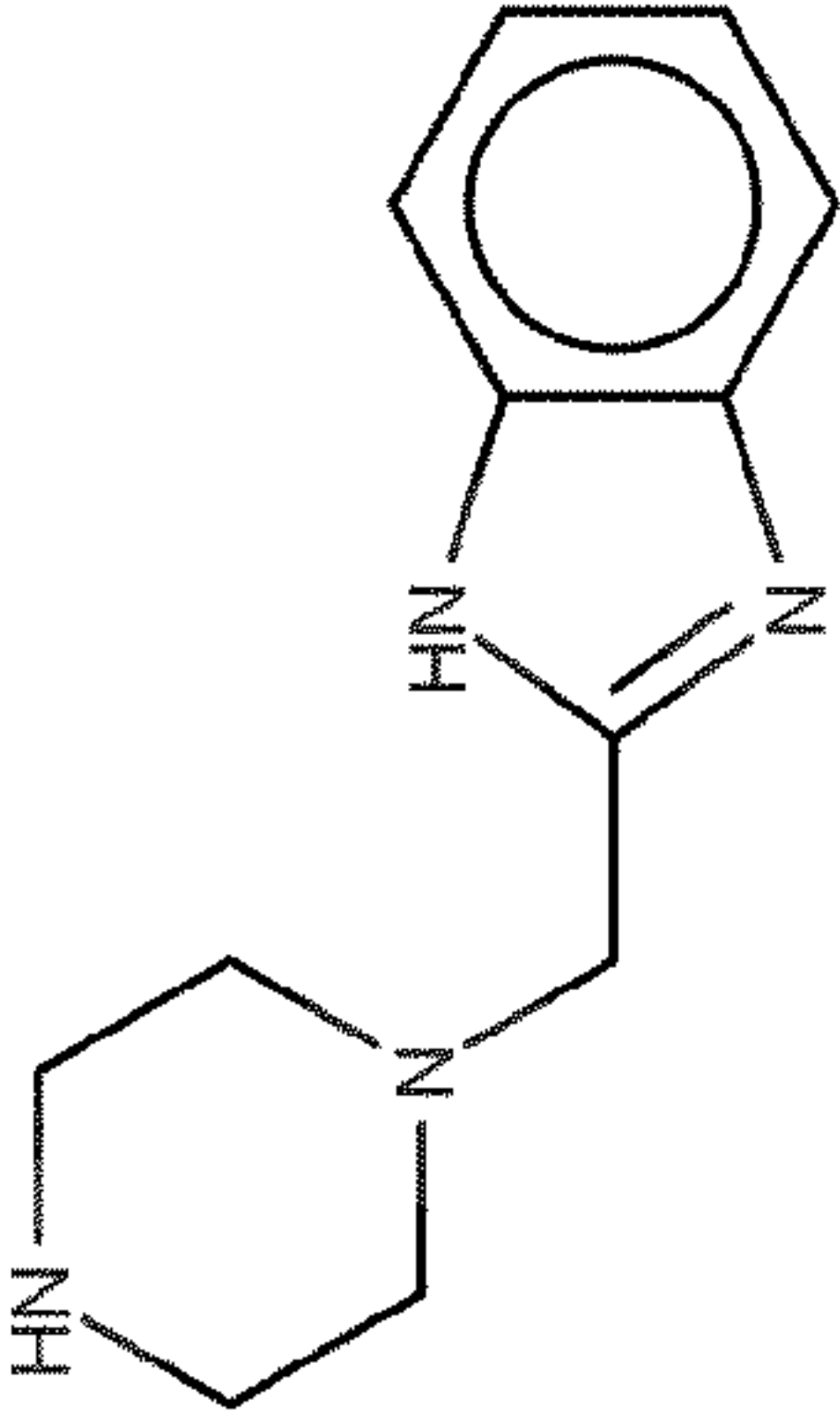
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 93		F2101-0054		F2101-0054_8_1 (1 of 18)
Compound 94		F2101-0132		F2101-0132_7_1 (1 of 18)
Compound 95		F2137-0016		F2137-0016_1_1 (1 of 4)

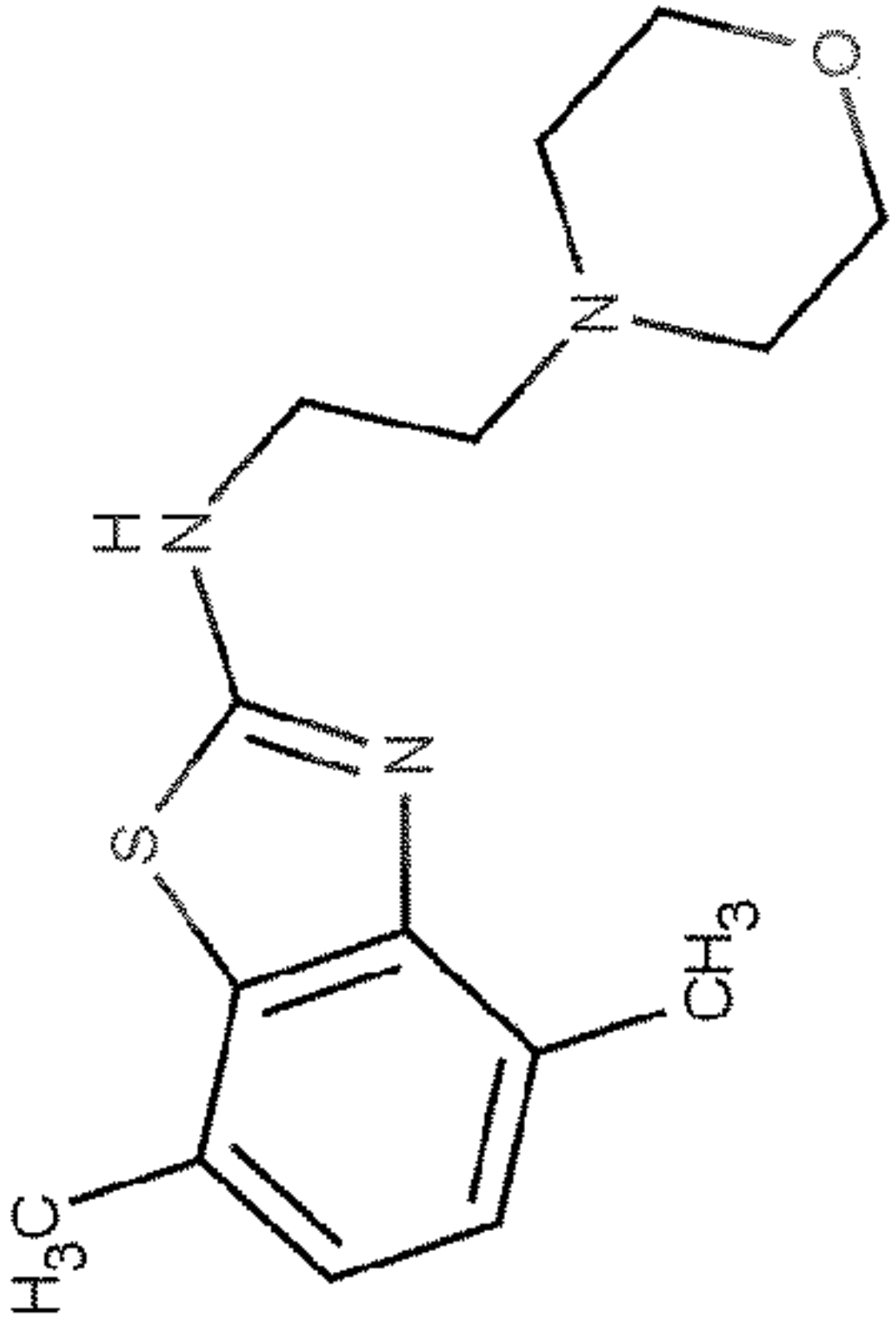
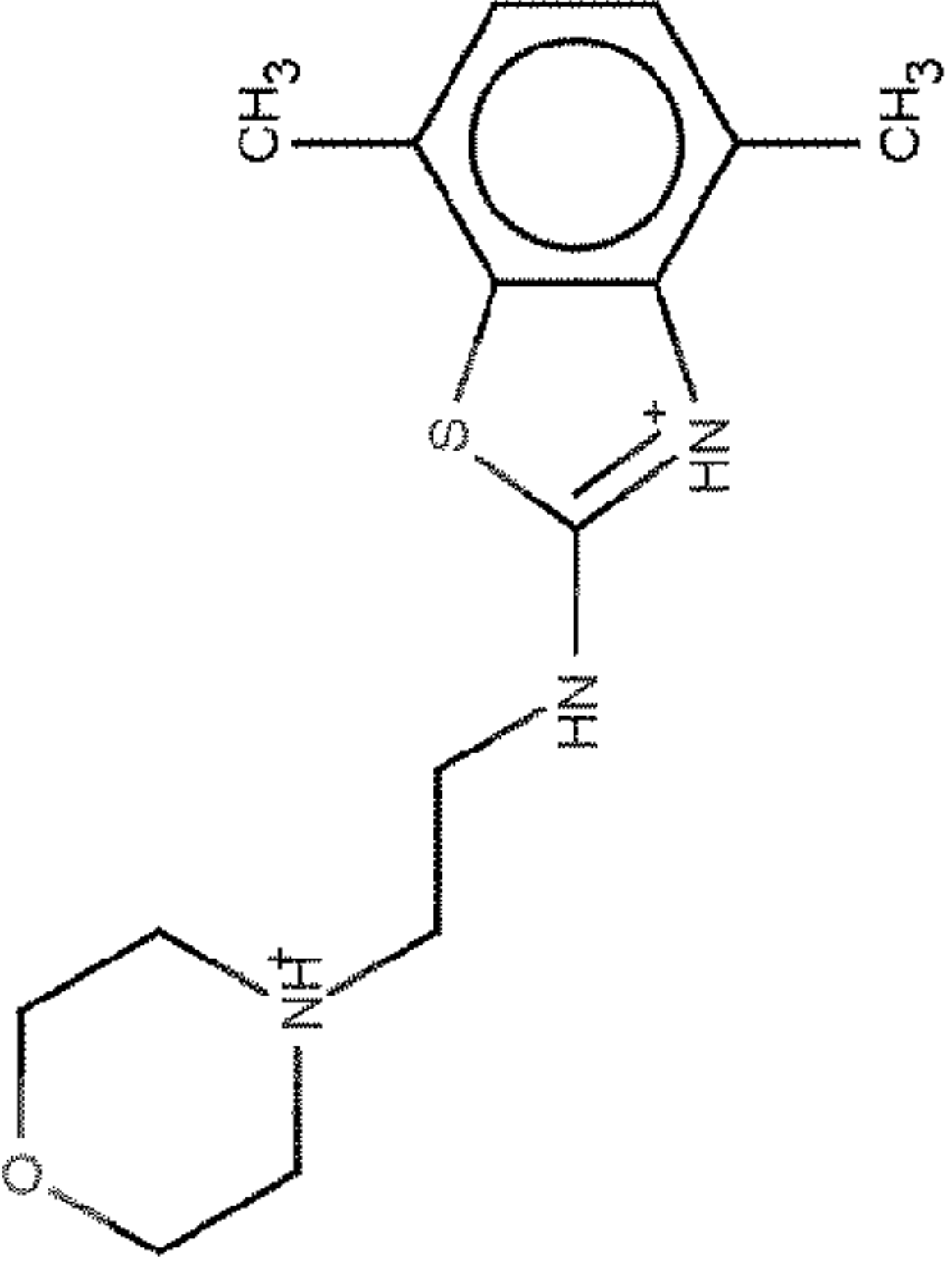
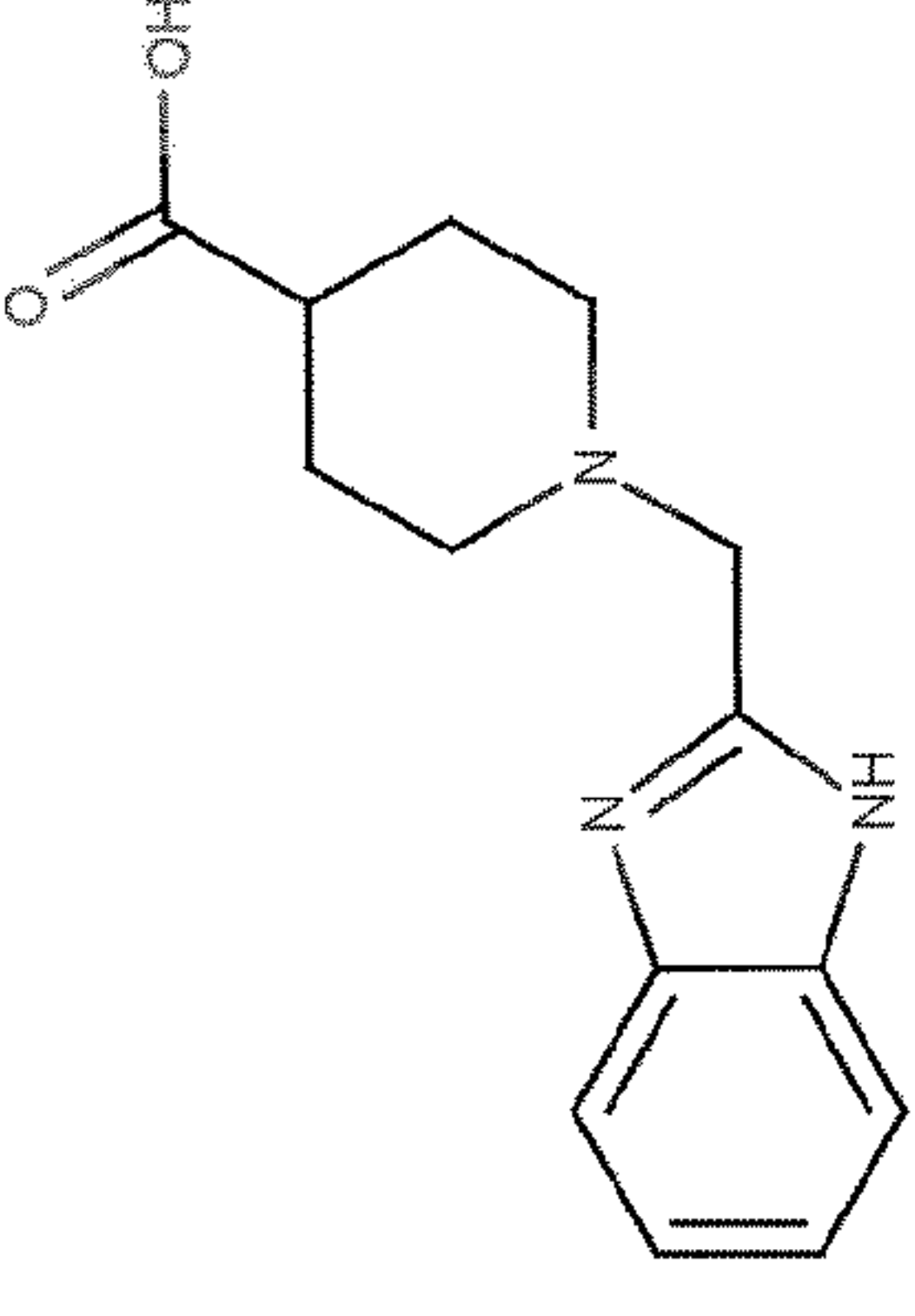
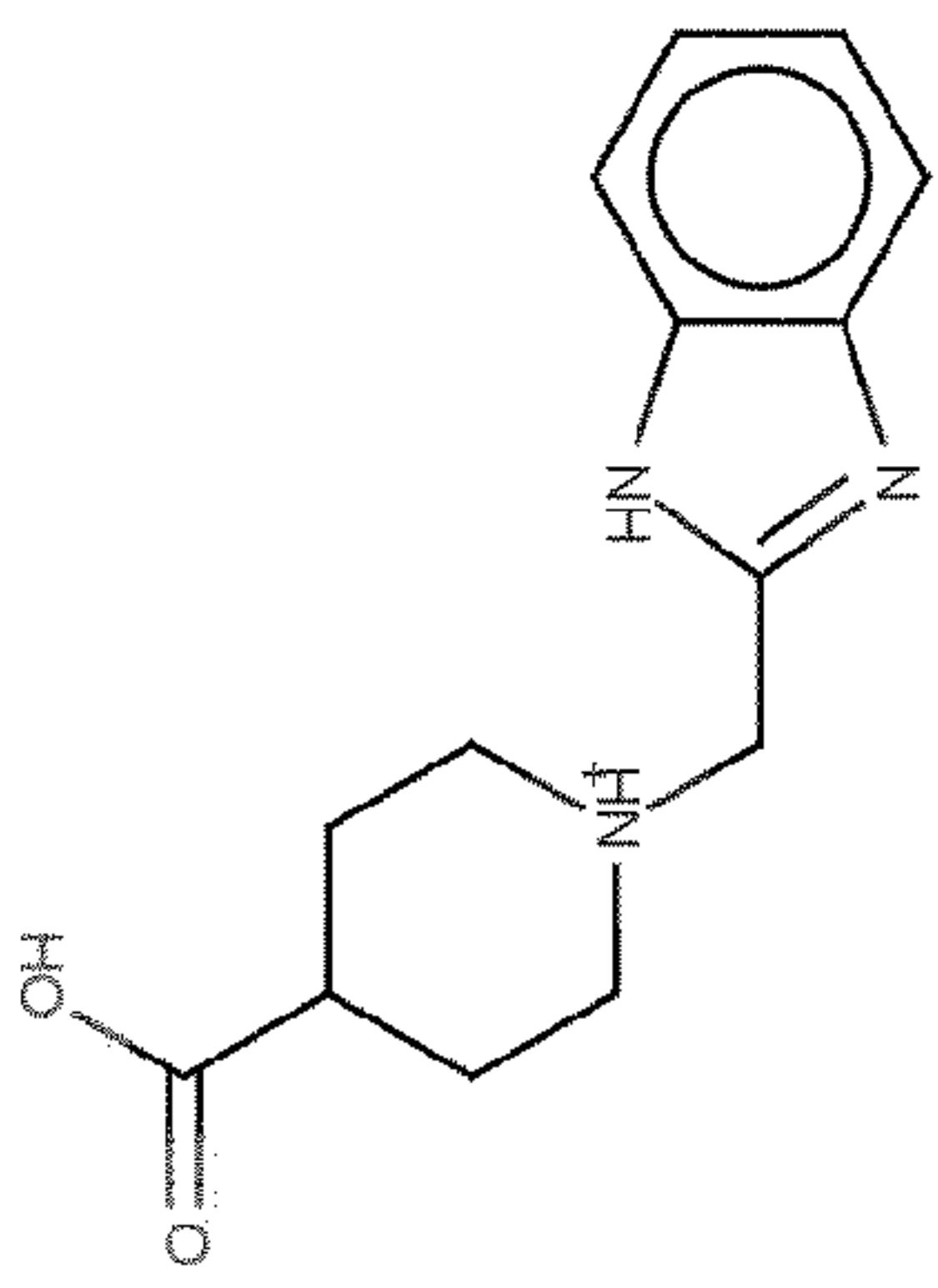
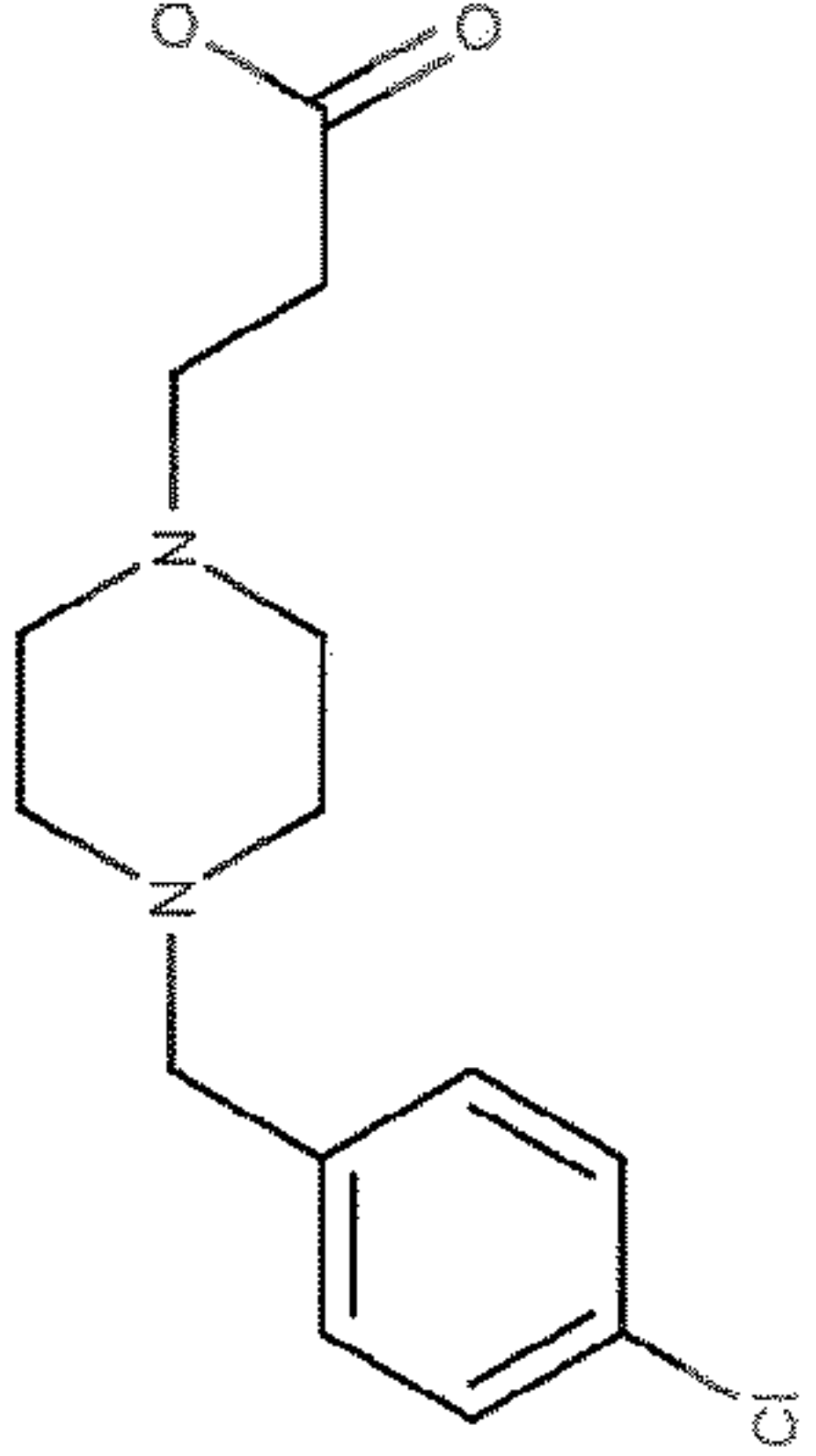
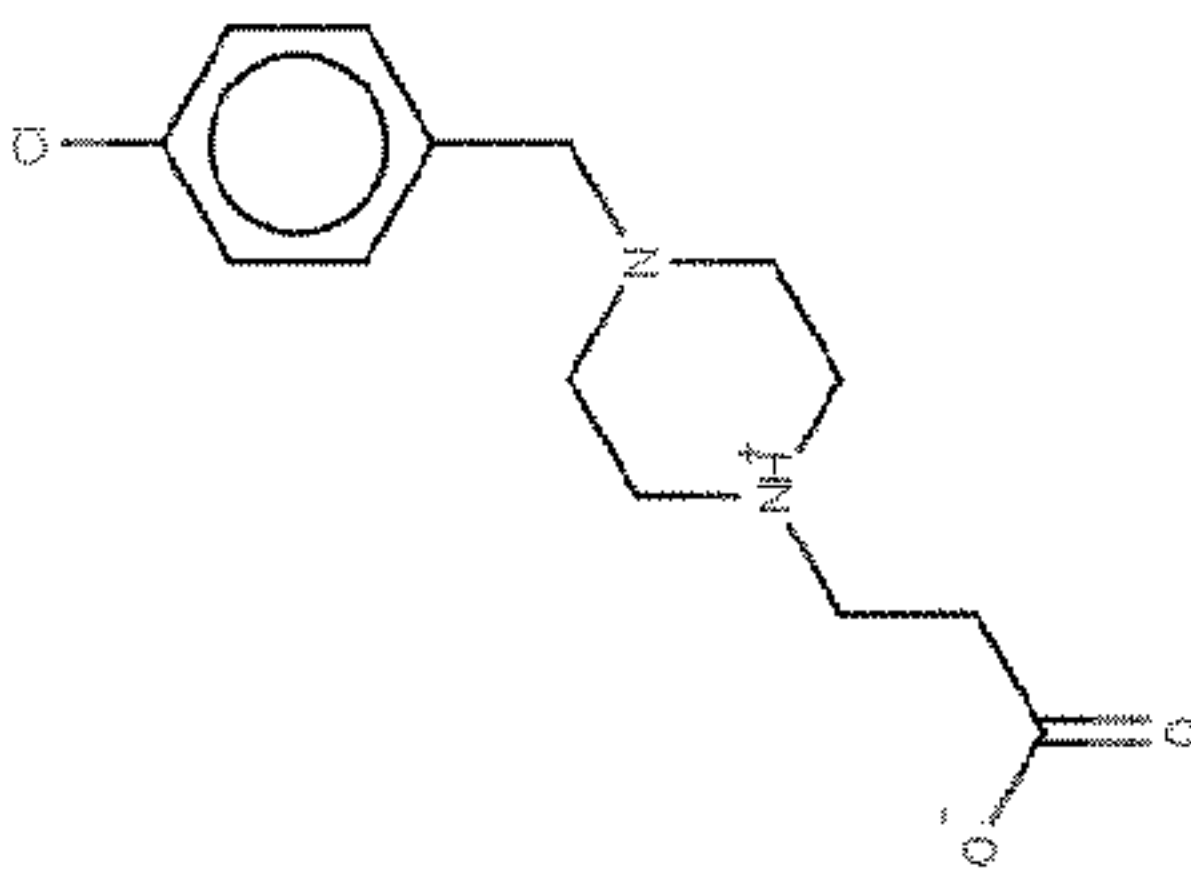
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 96		F2145-0090		F2145-0090_3_1 (1 of 4)
Compound 97		F2145-0296		F2145-0296_3_1 (1 of 4)
Compound 98		F2145-0373		F2145-0373_3_1 (1 of 7)

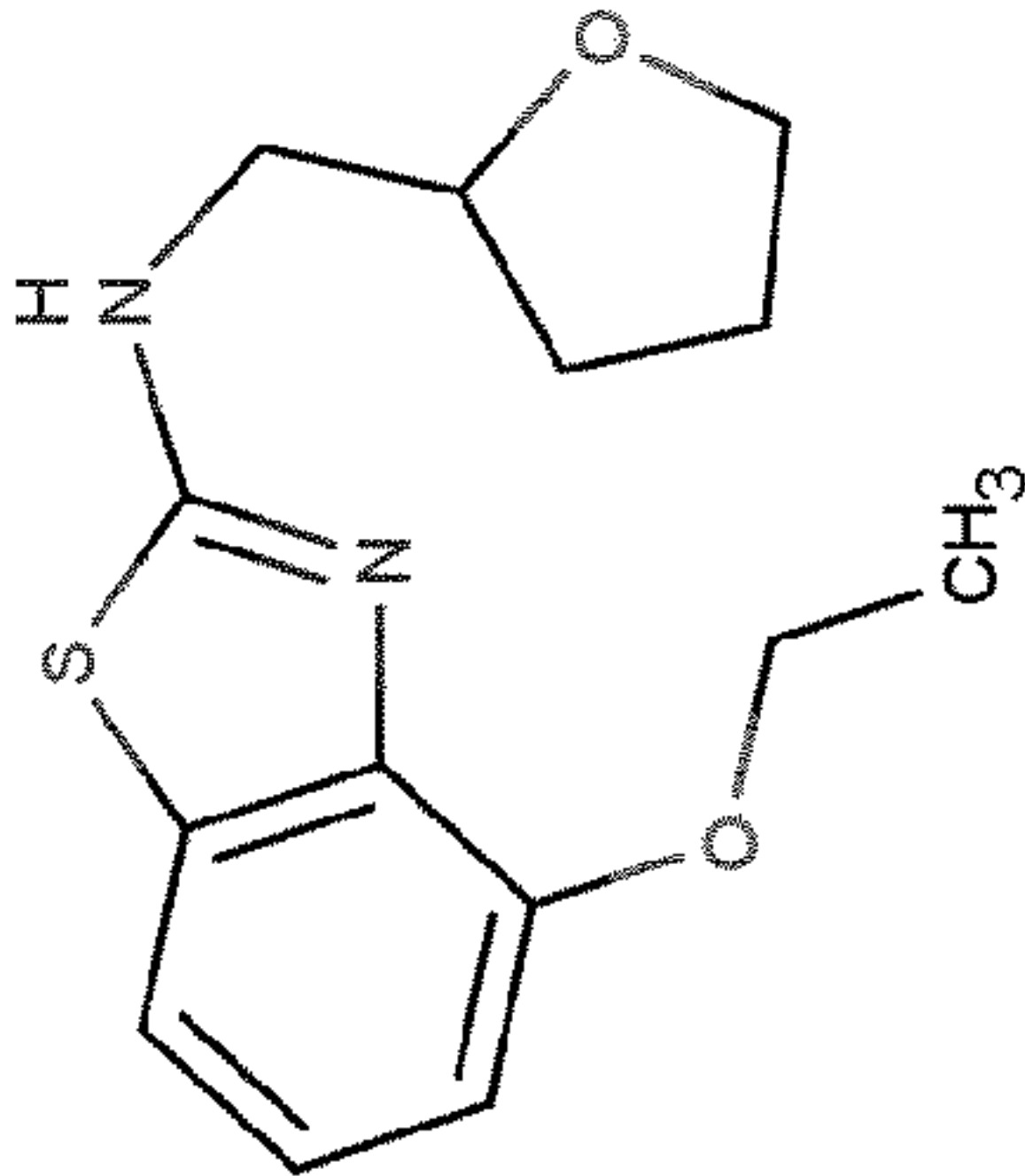
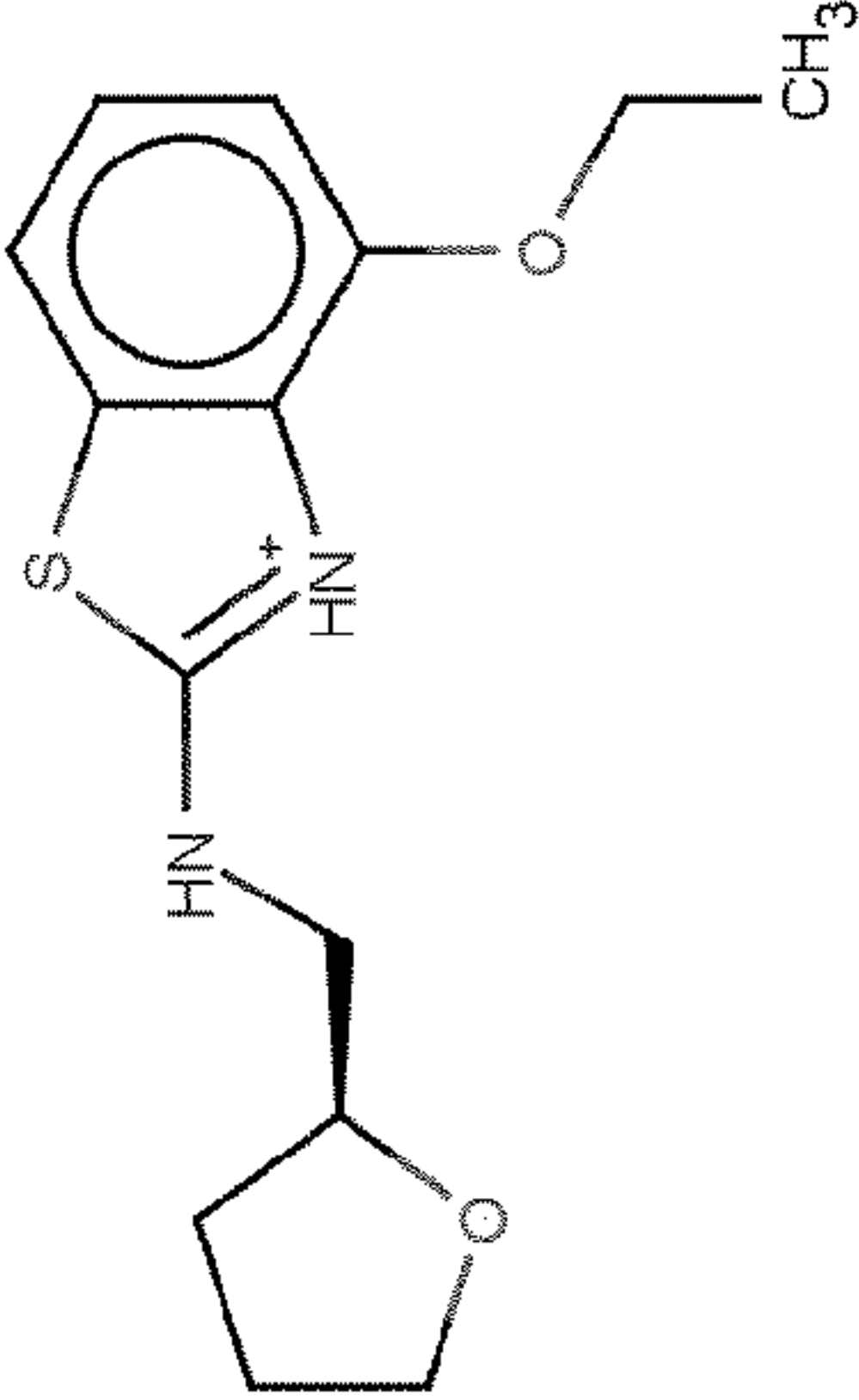
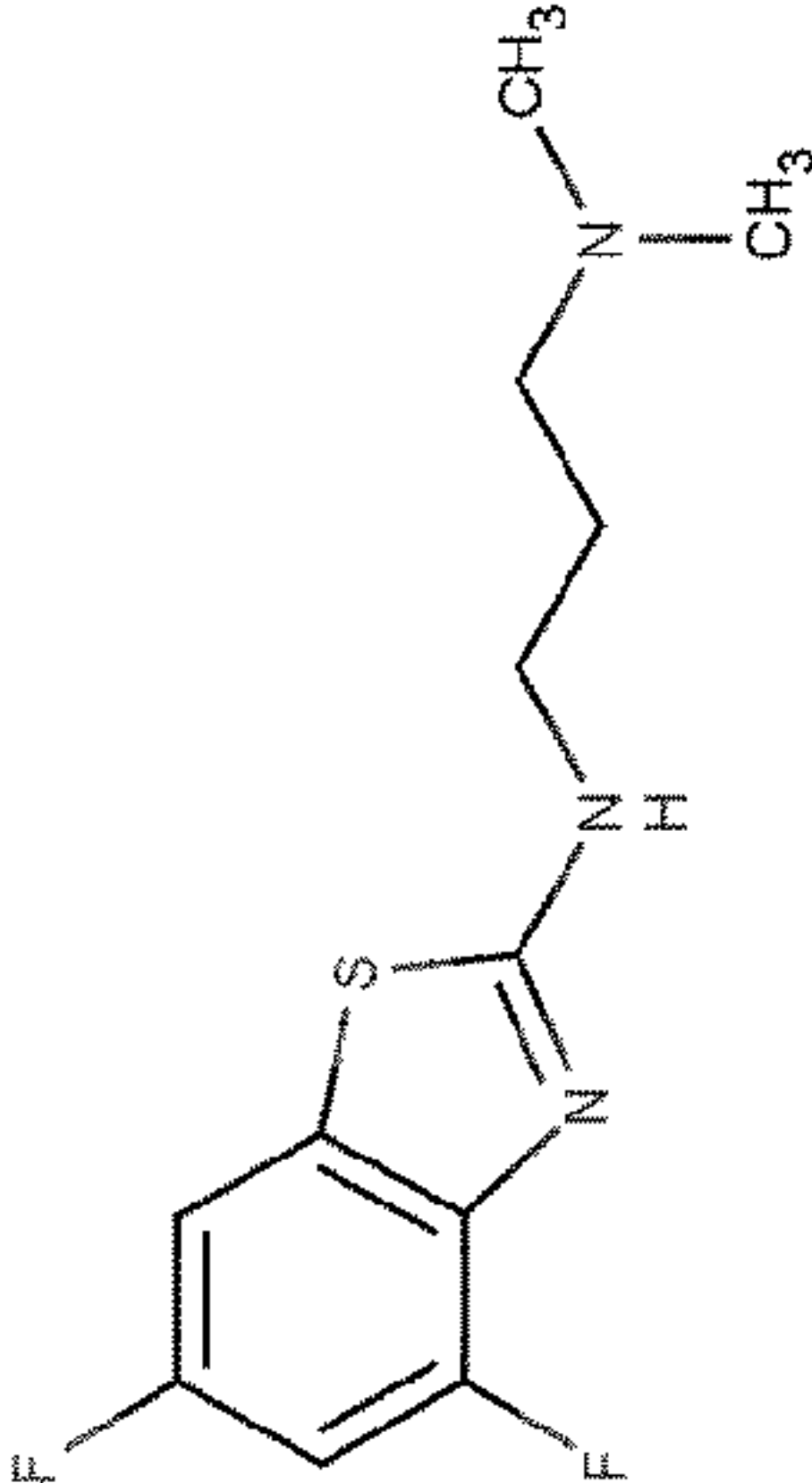
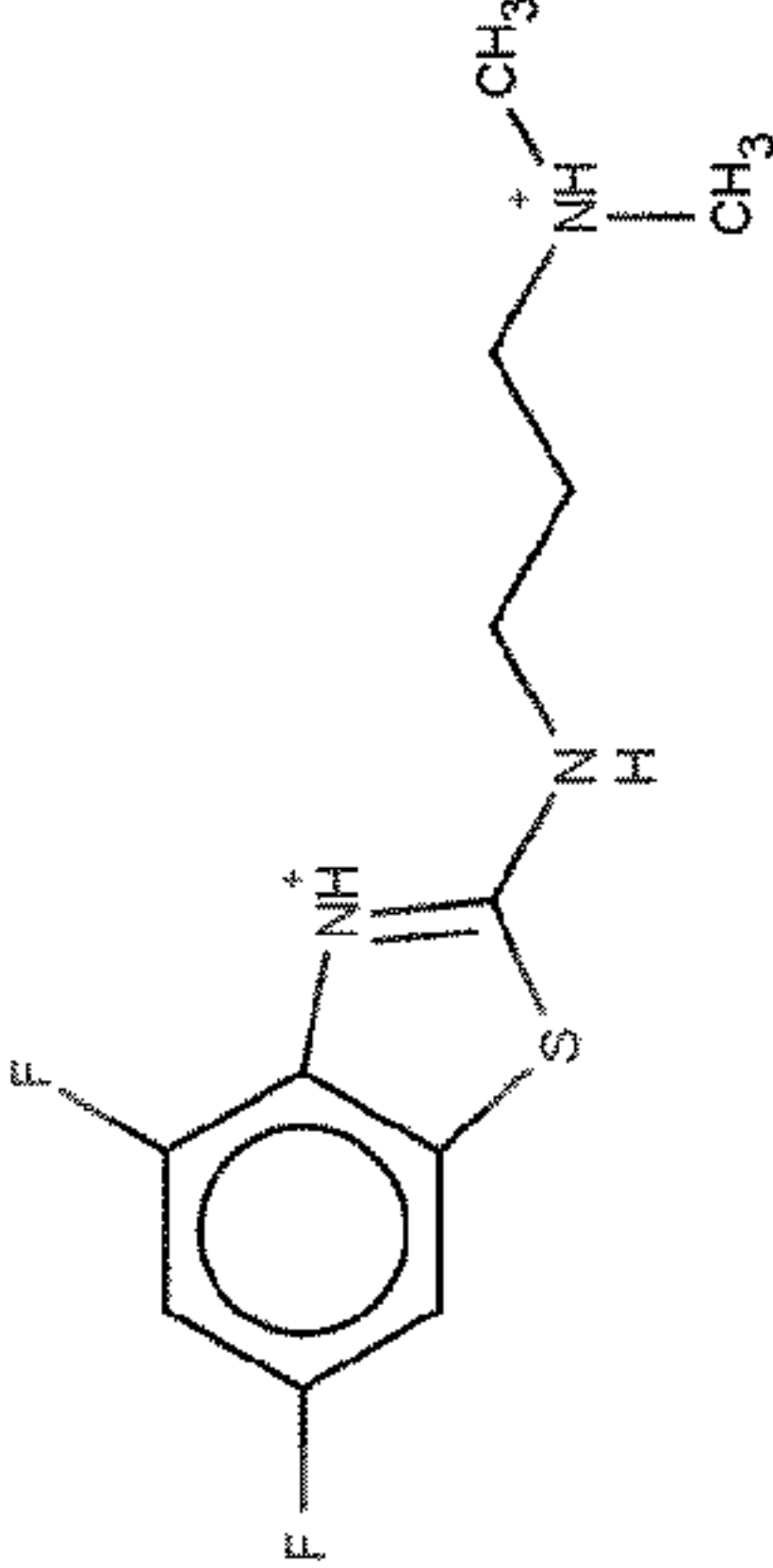
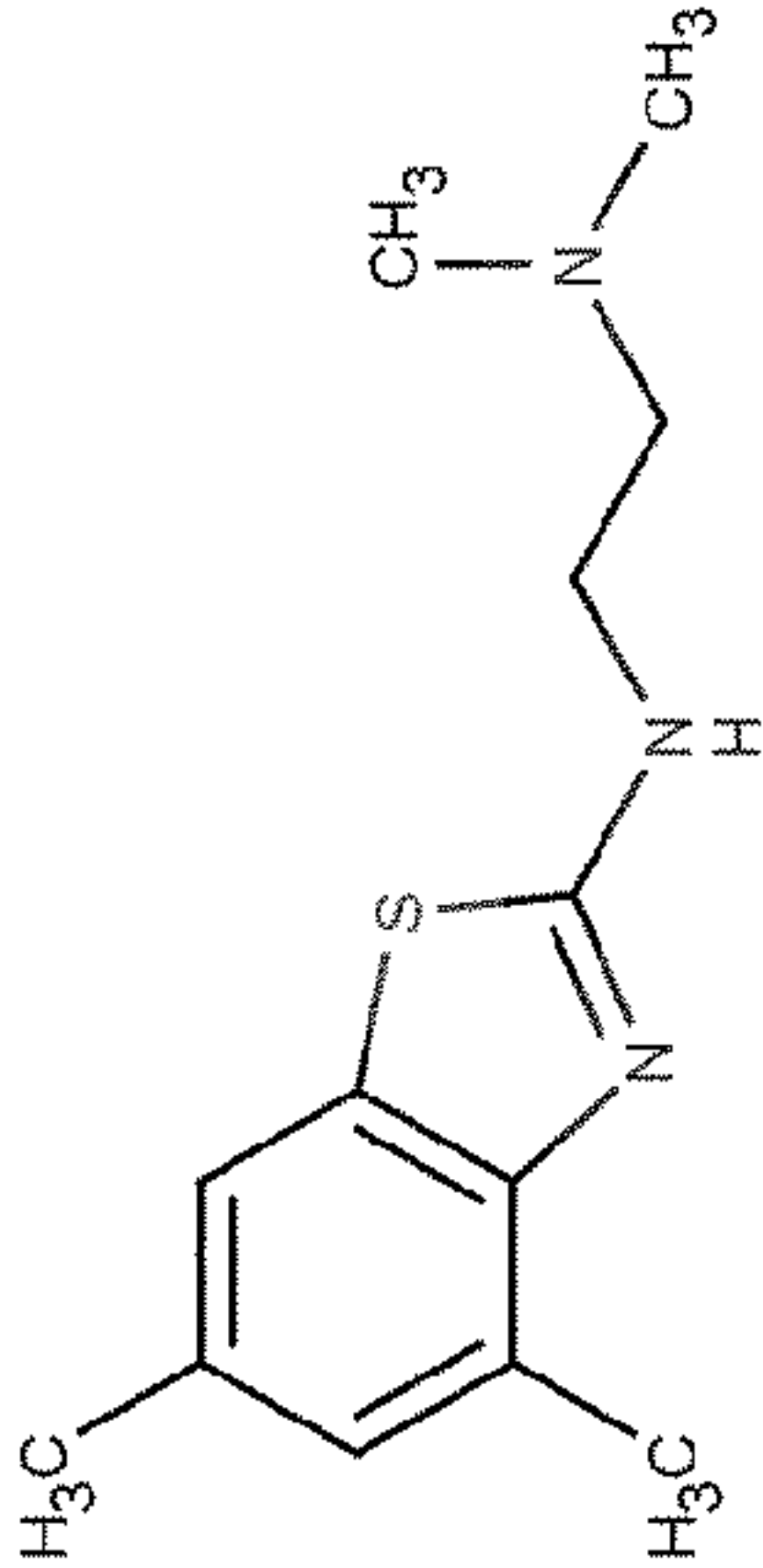
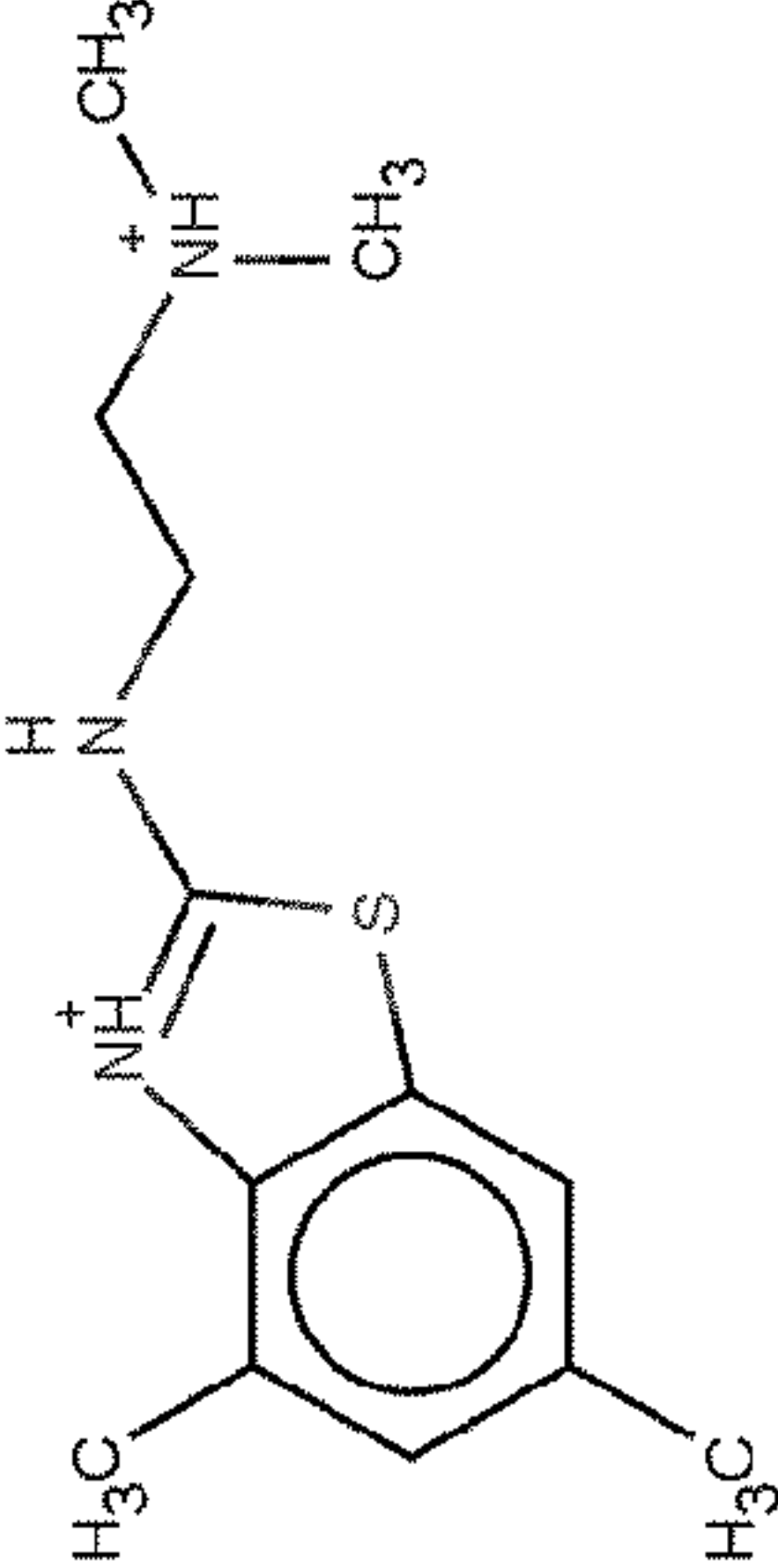
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 99		F2145-0423		F2145-0423_5_1 (1 of 7)
Compound 100		F2145-0424		F2145-0424_1_1 (1 of 7)
Compound 102		F2145-0478		F2145-0478_2_1 (1 of 4)

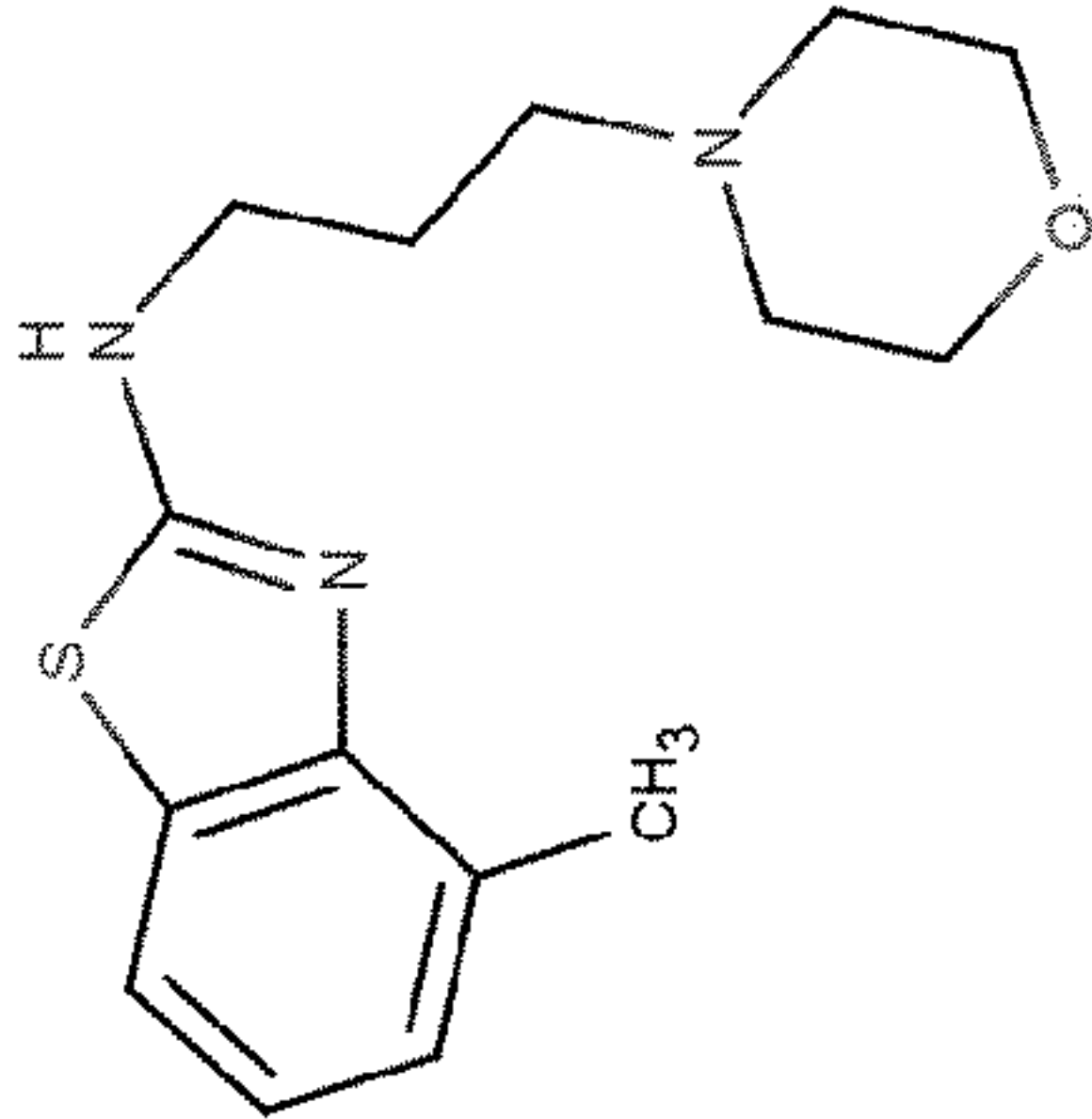
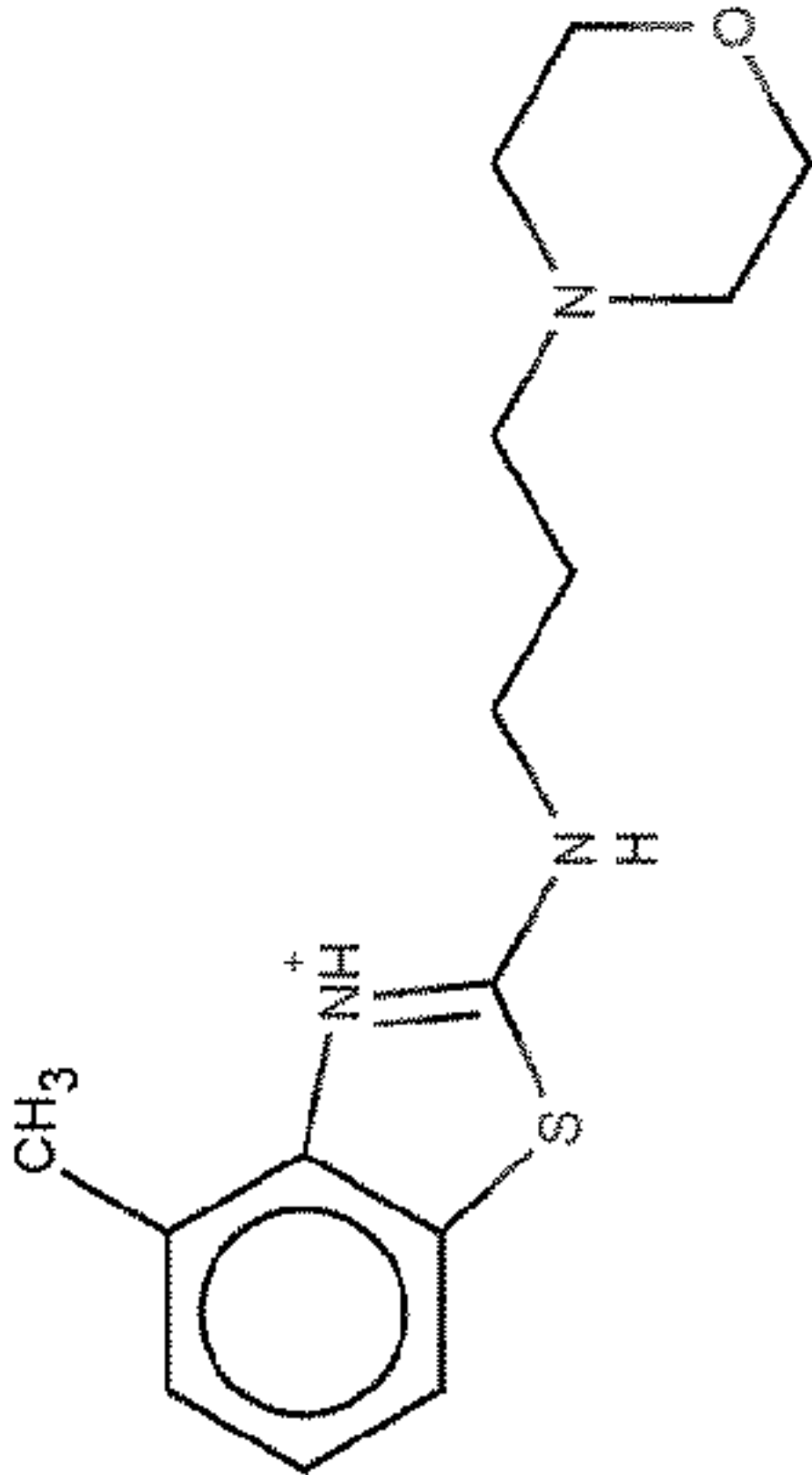
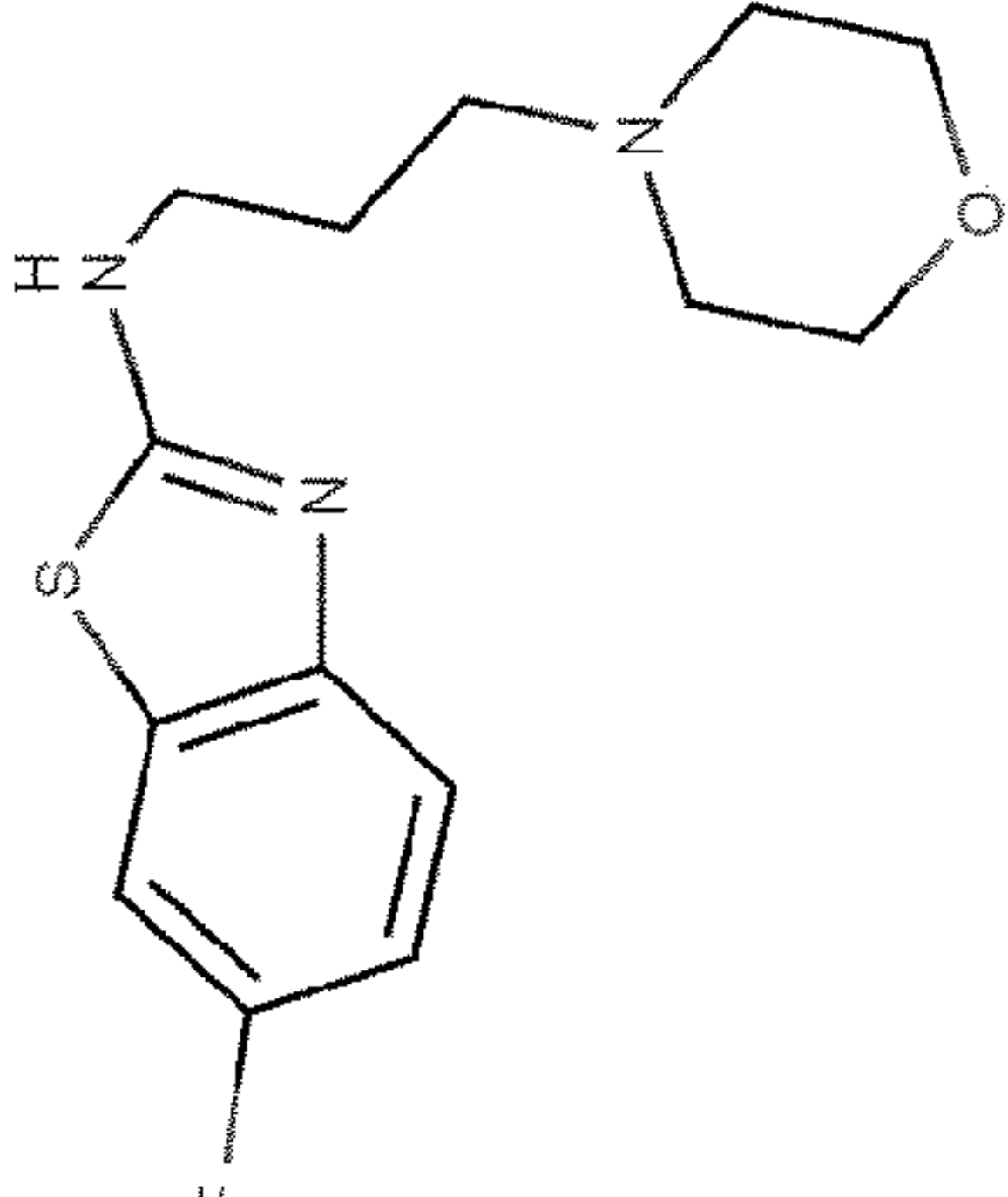
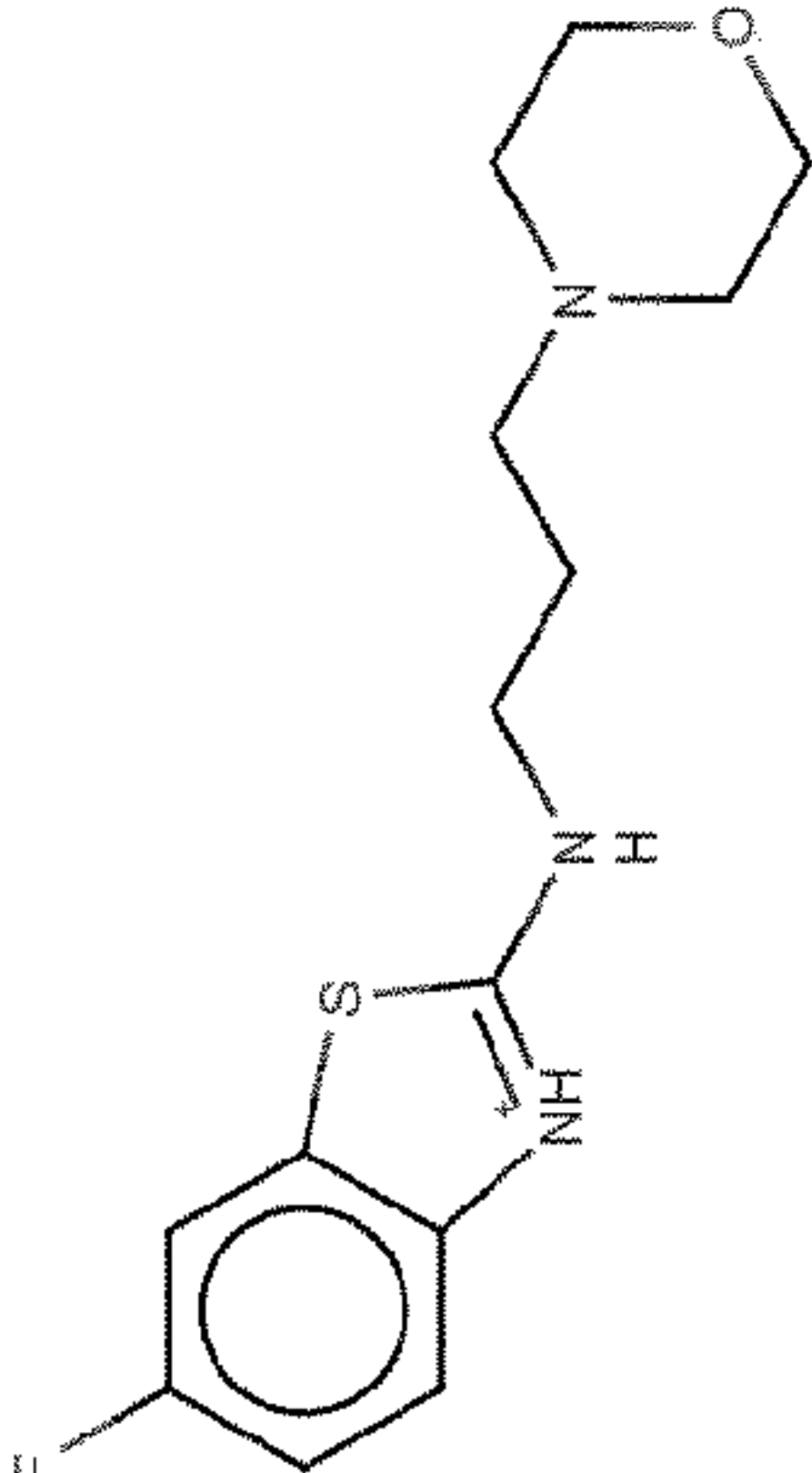
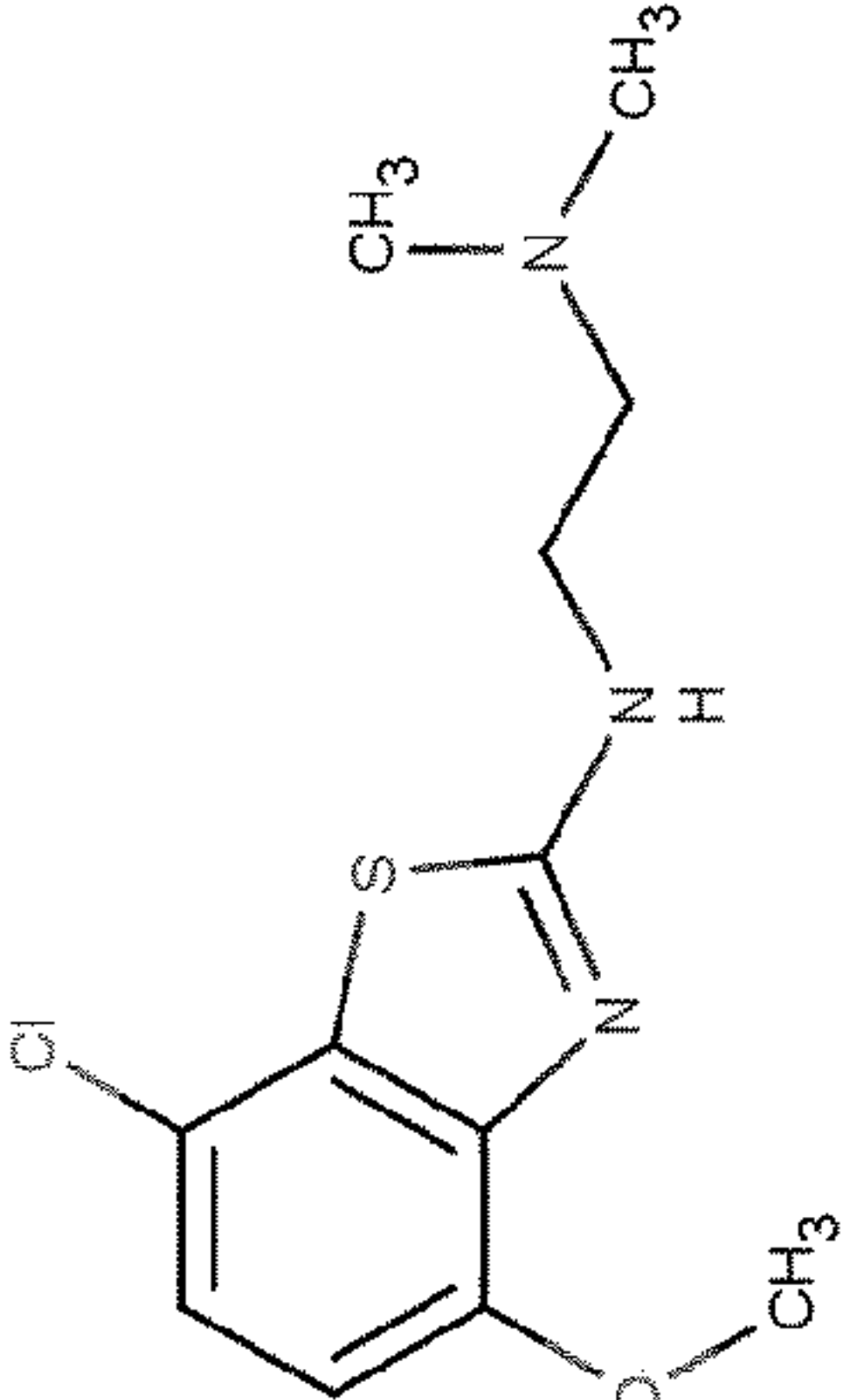
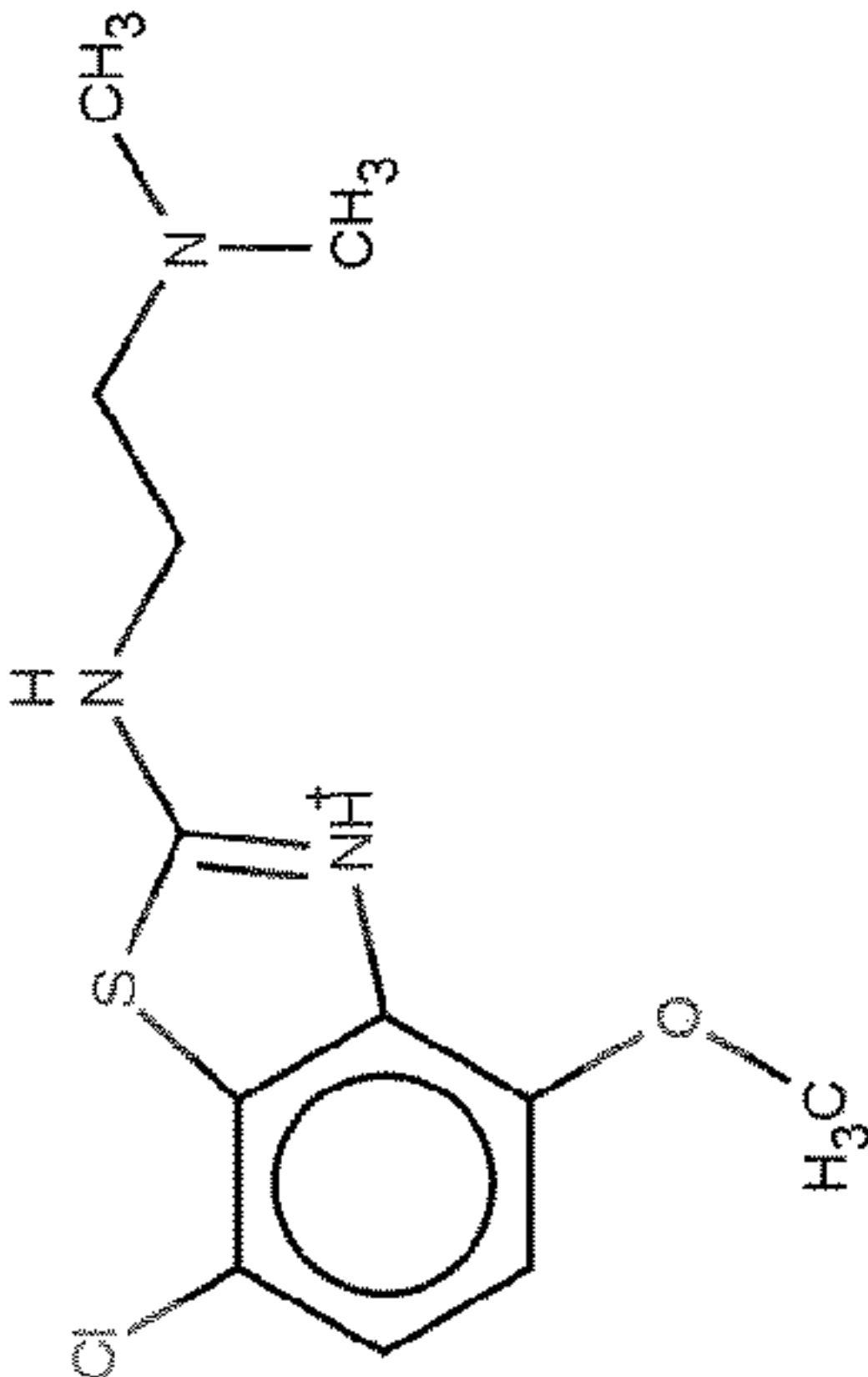
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 103		F2146-0582		F2146-0582_3_1 (1 of 4)
Compound 104		F2147-0176		F2147-0176_1_1 (1 of 4)
Compound 105		F2147-0549		F2147-0549_3_2 (1 of 8)

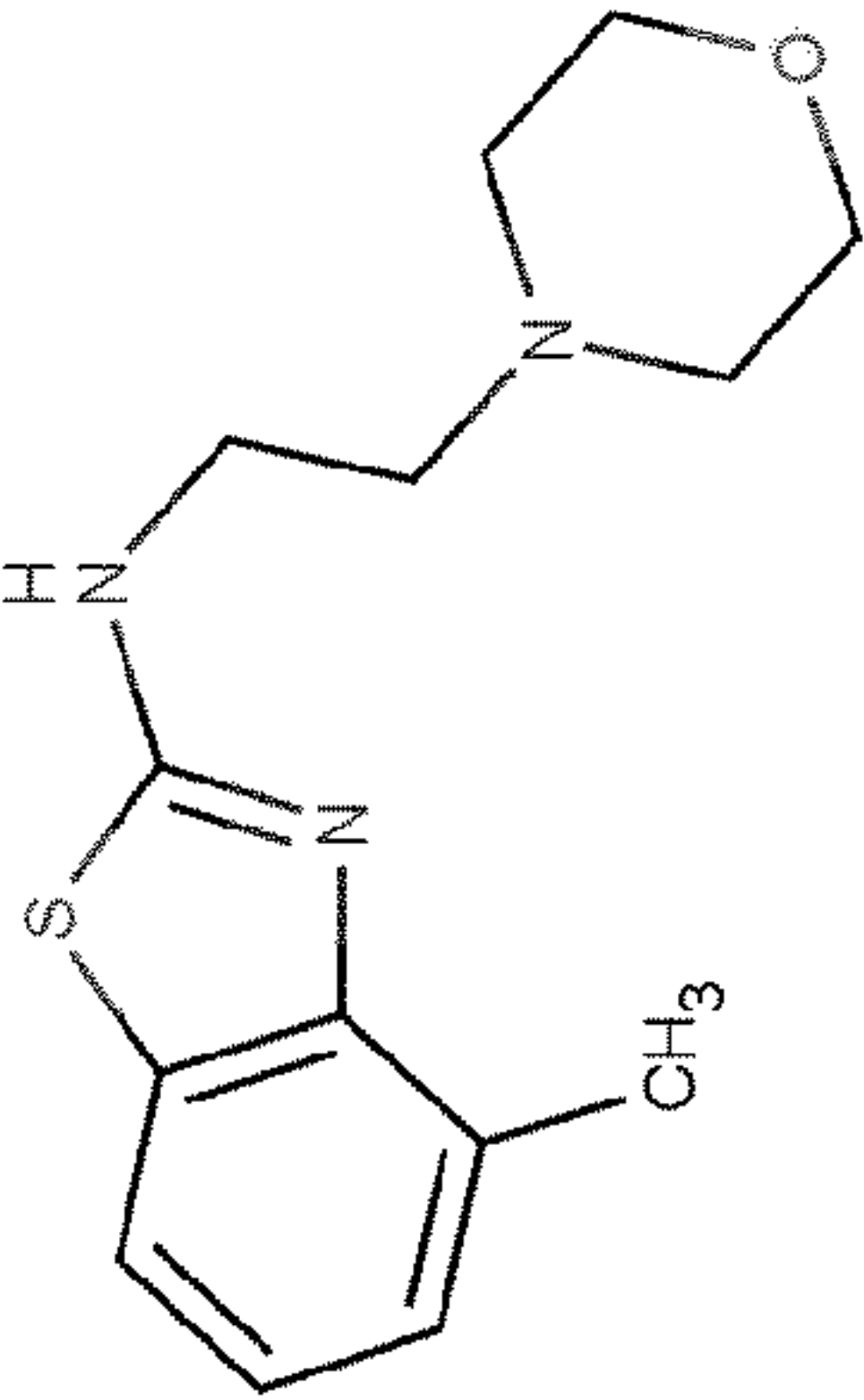
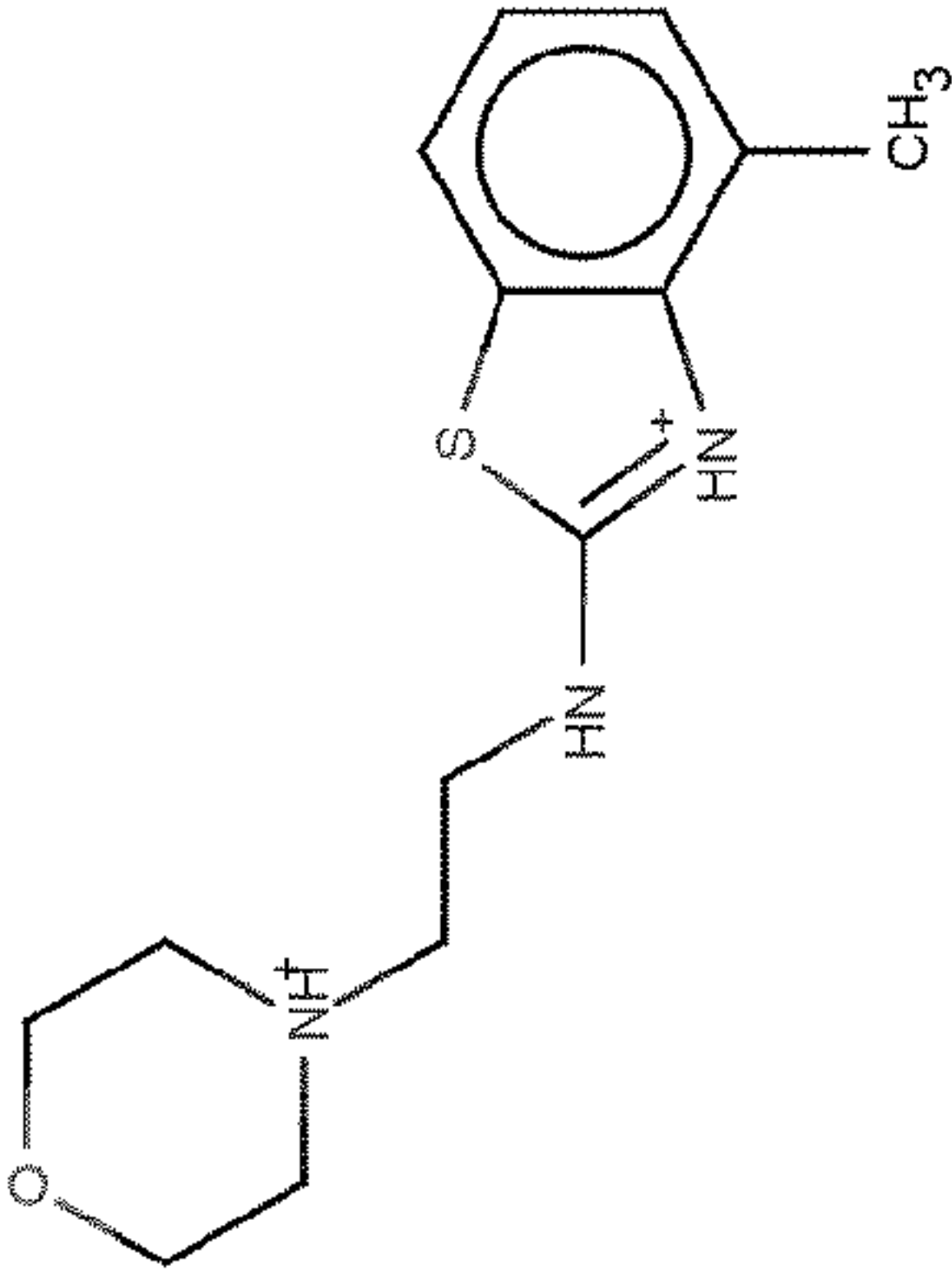
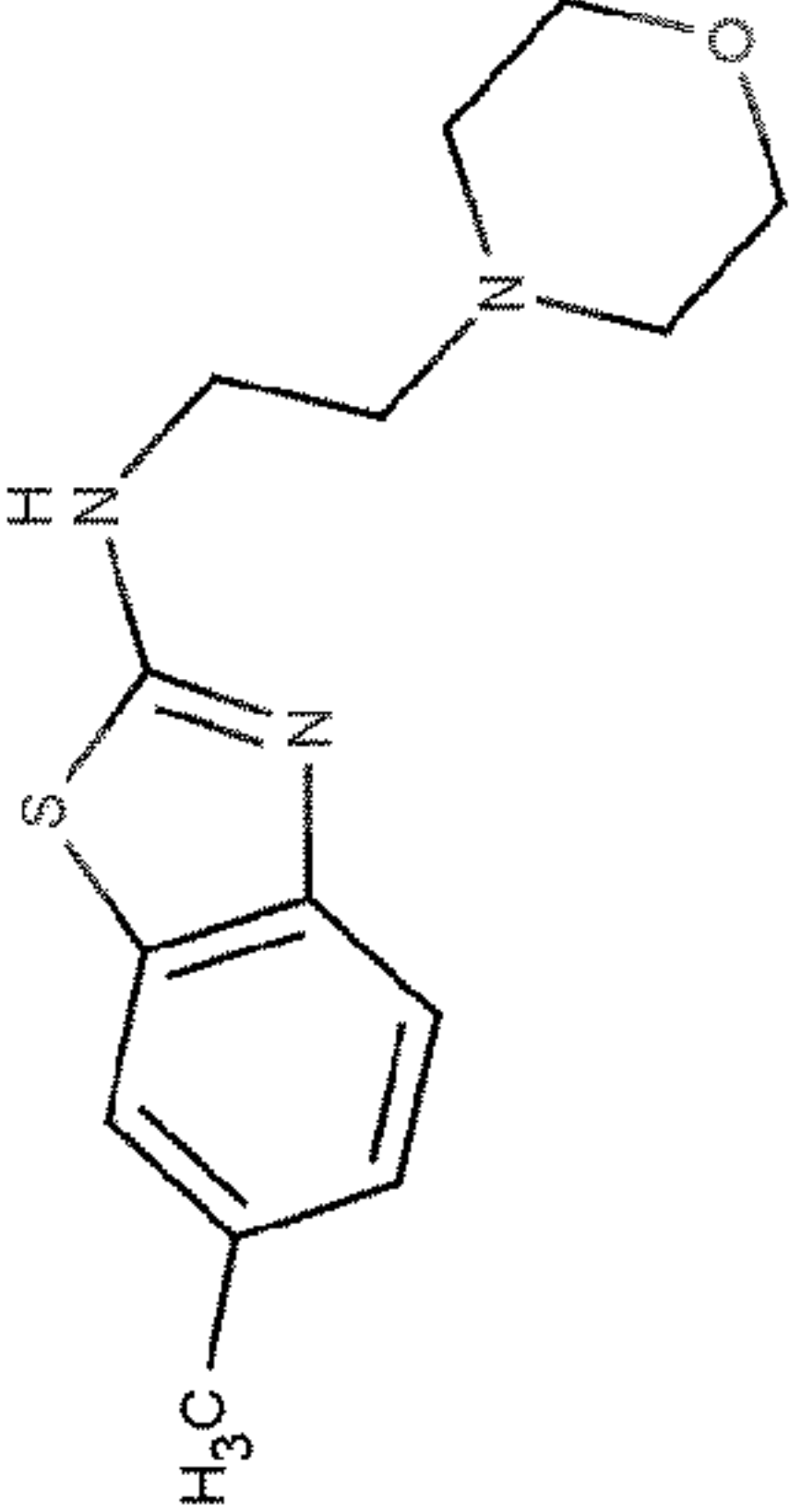
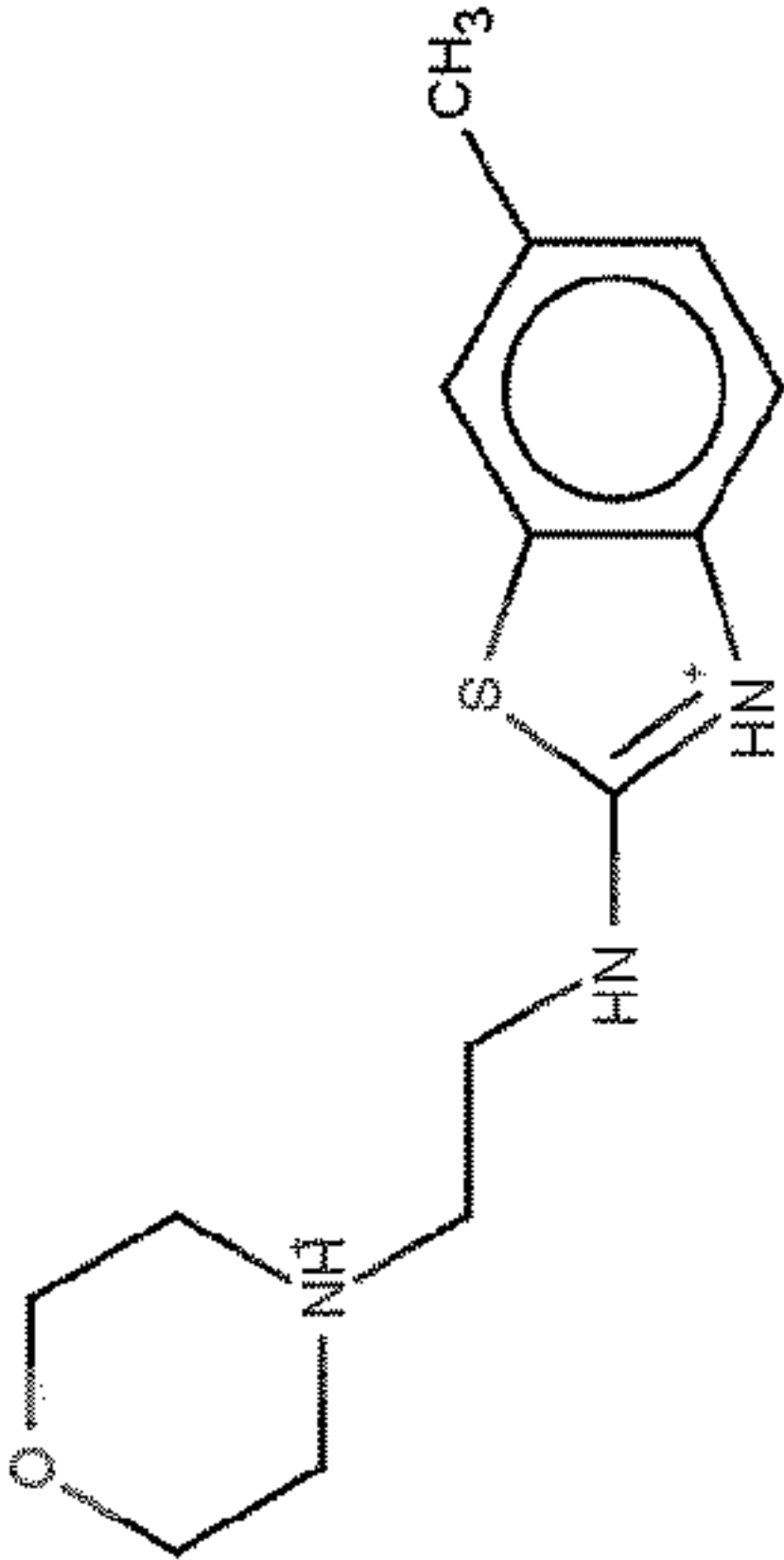
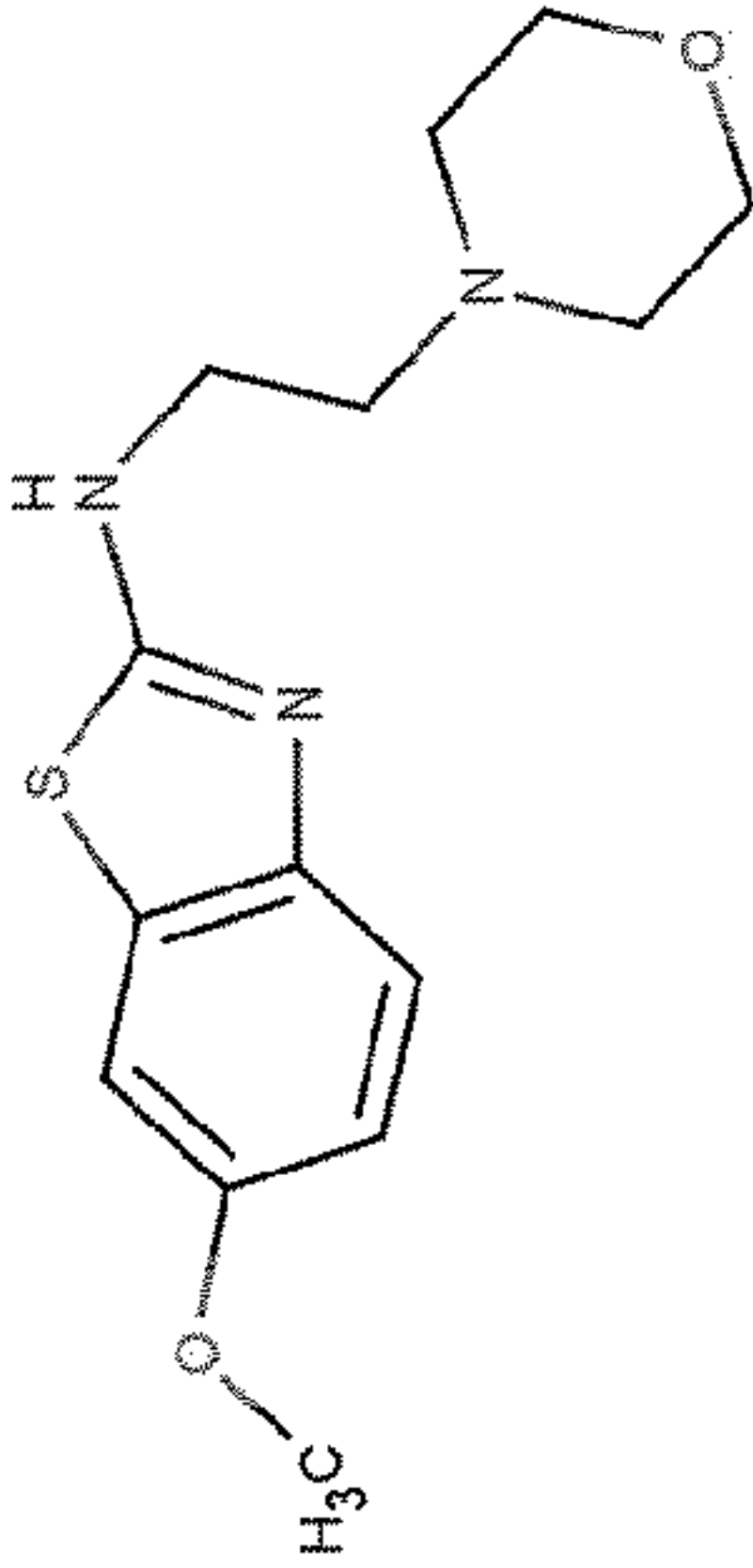
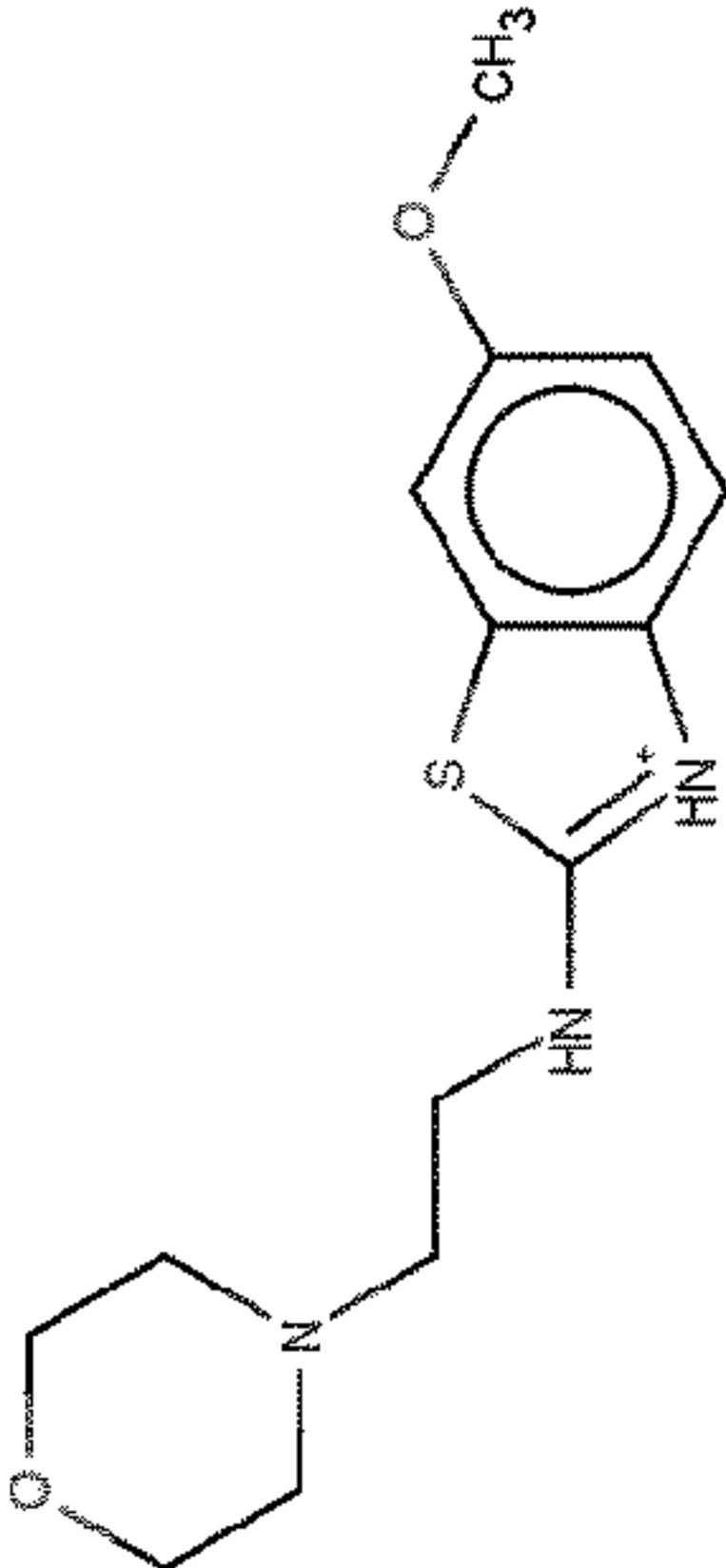
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 106		F2148-0005		F2148-0005_6_4 (1 of 28)
Compound 107		F2148-0237		F2148-0237_2_1 (1 of 7)
Compound 108		F2150-0003		F2150-0003_3_1 (1 of 4)

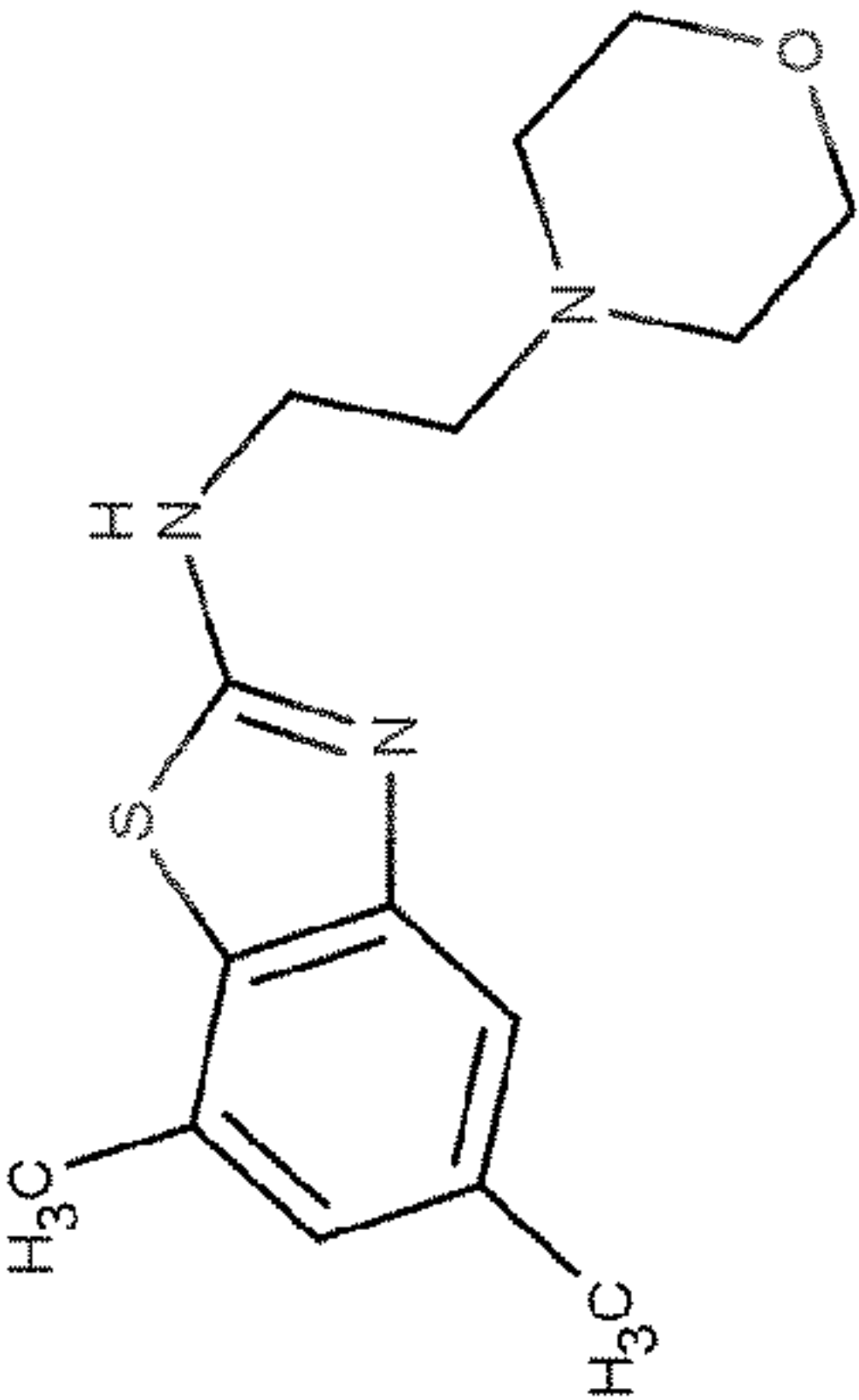
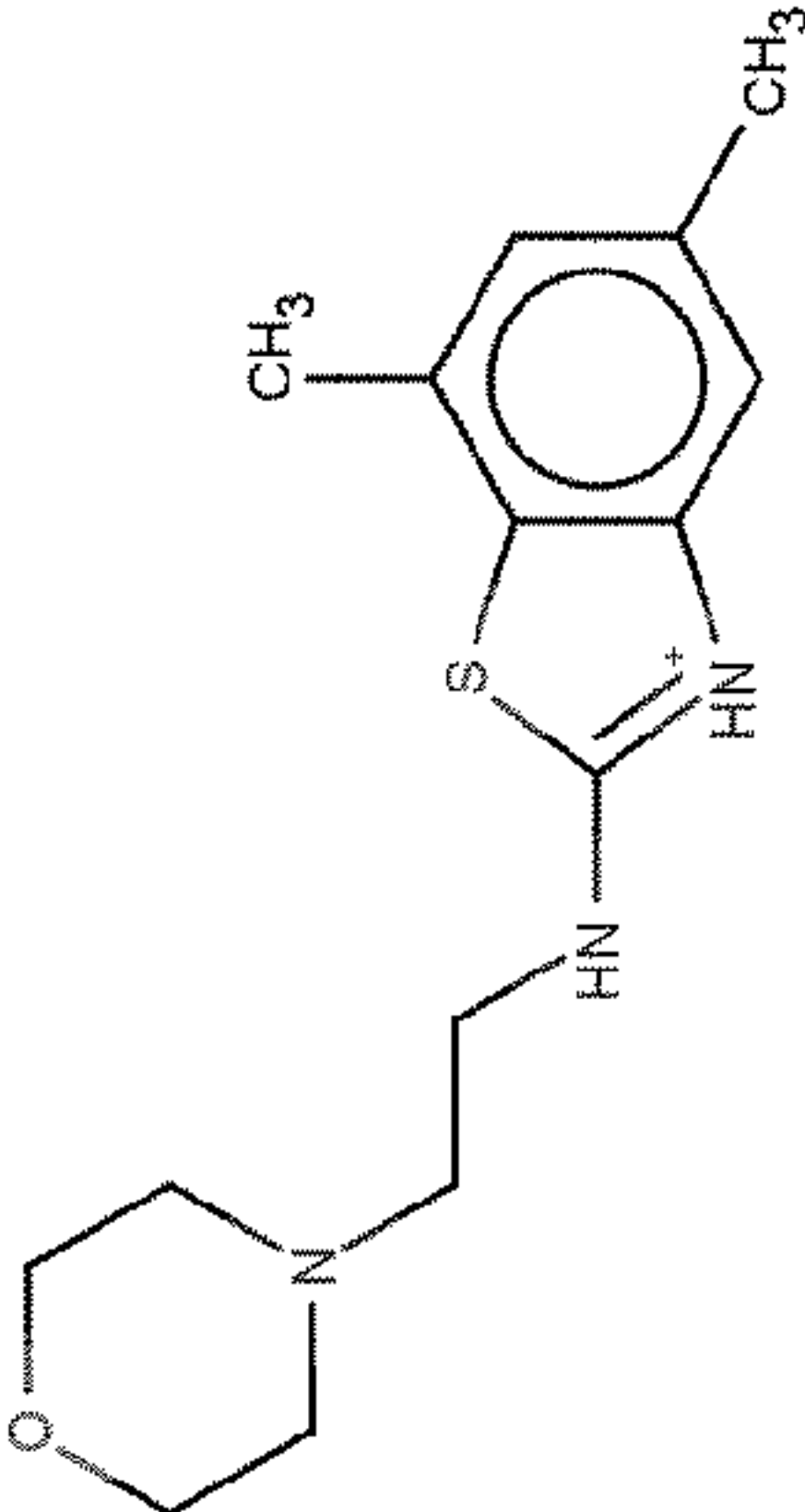
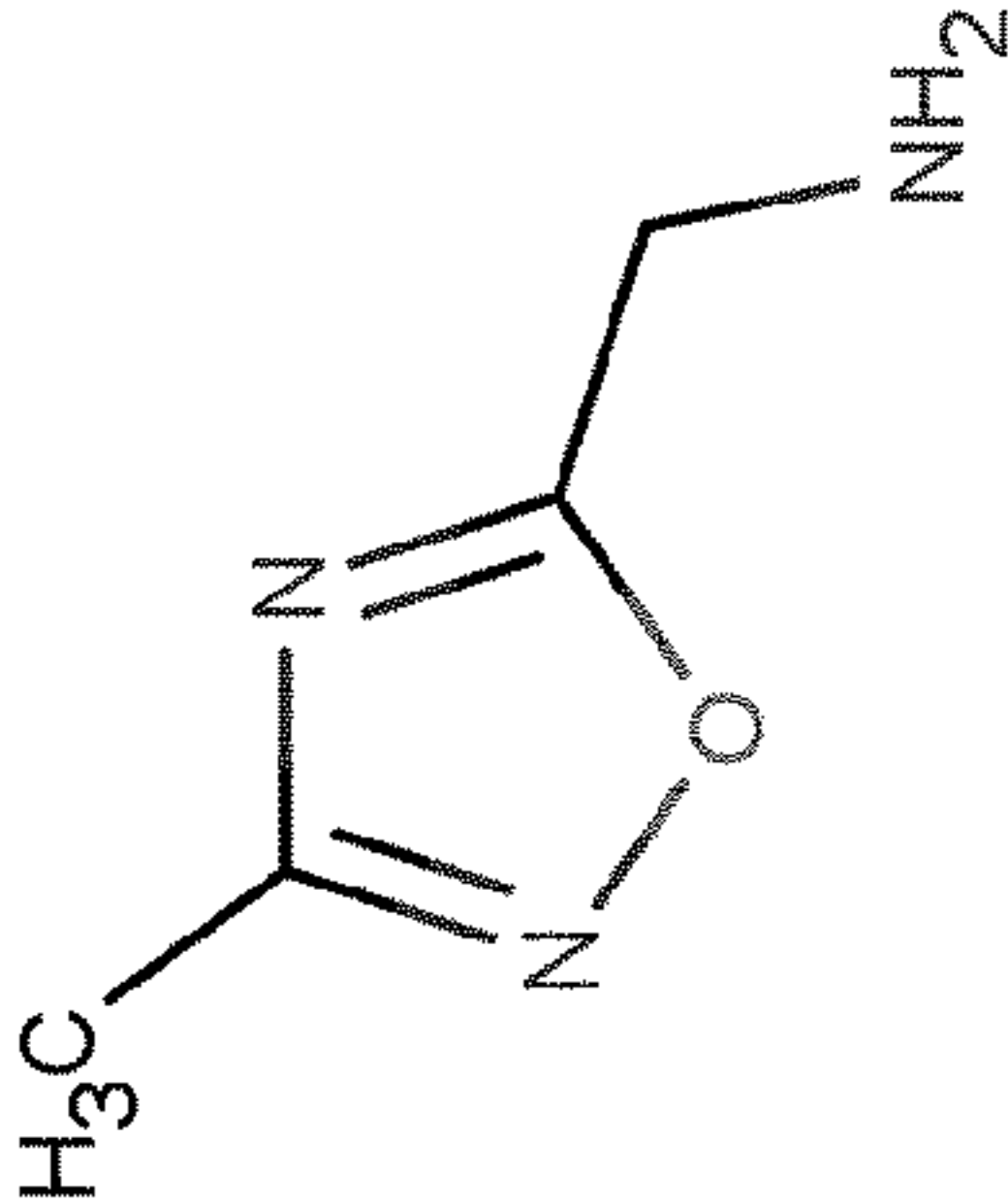
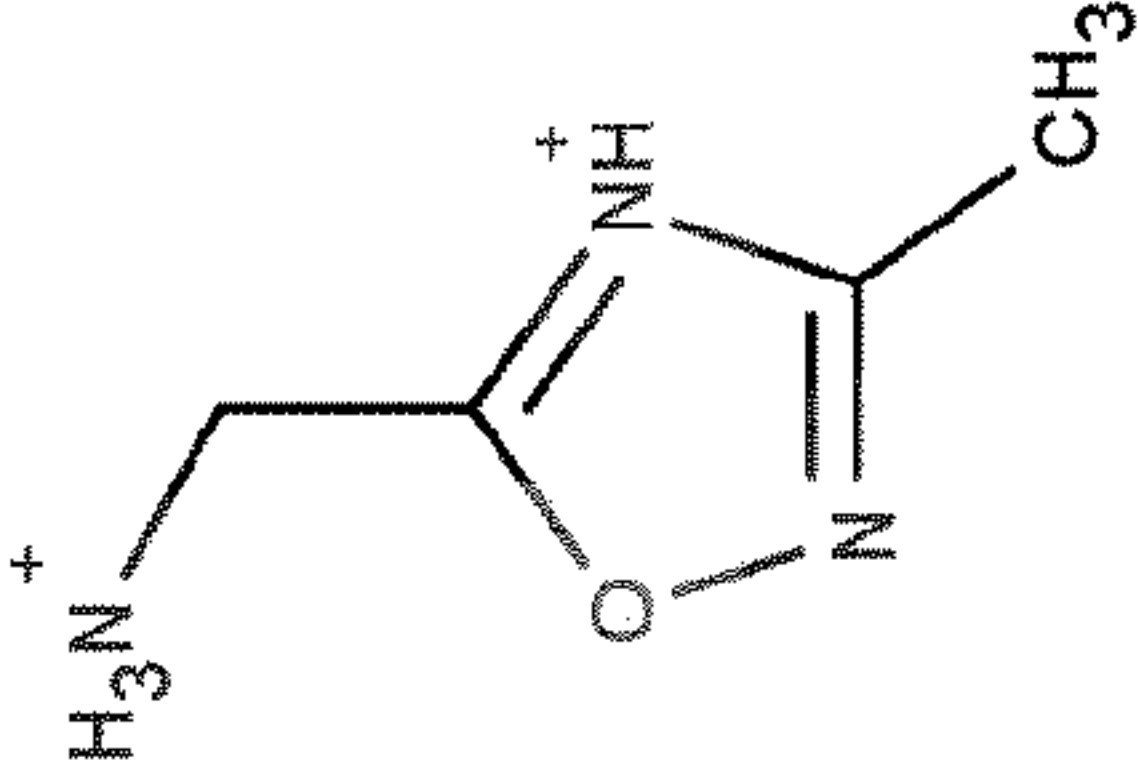
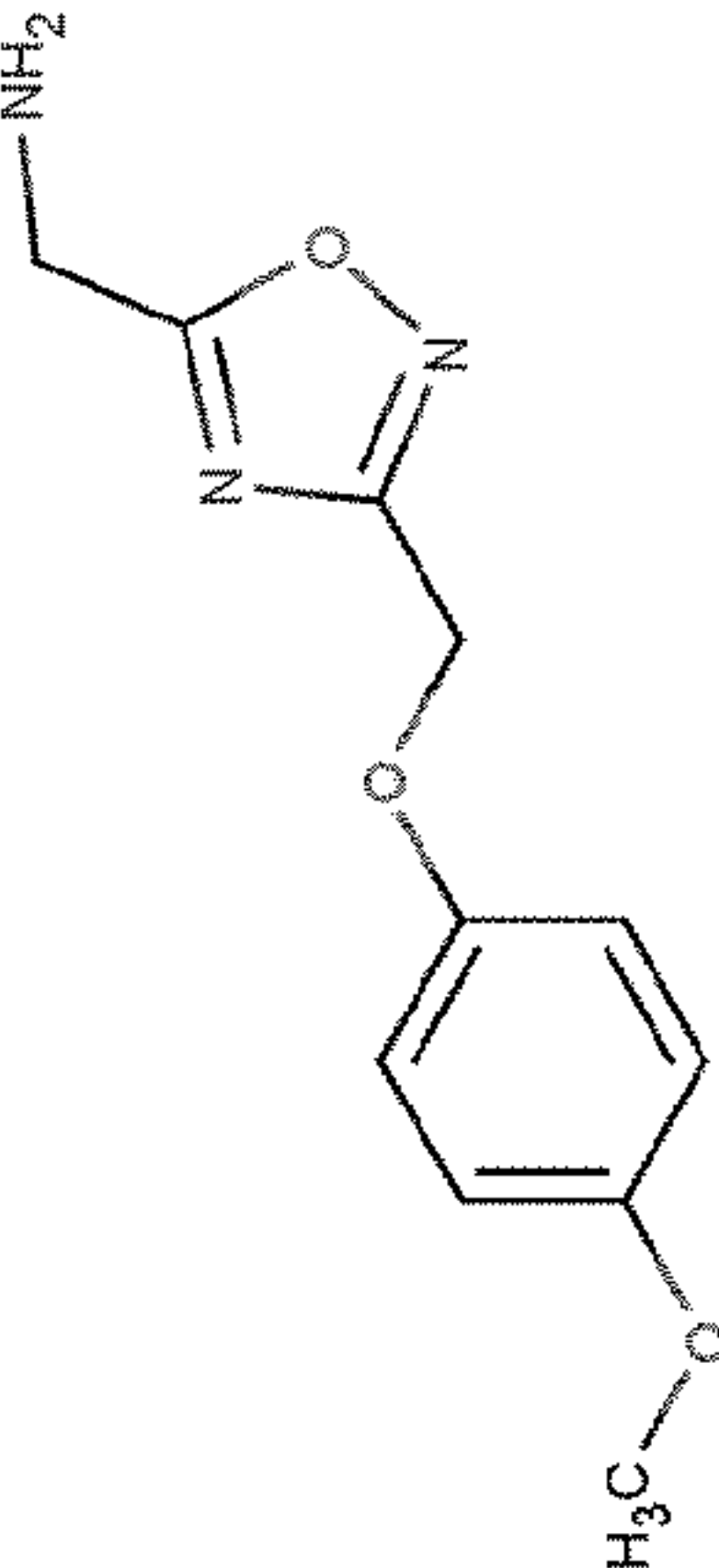
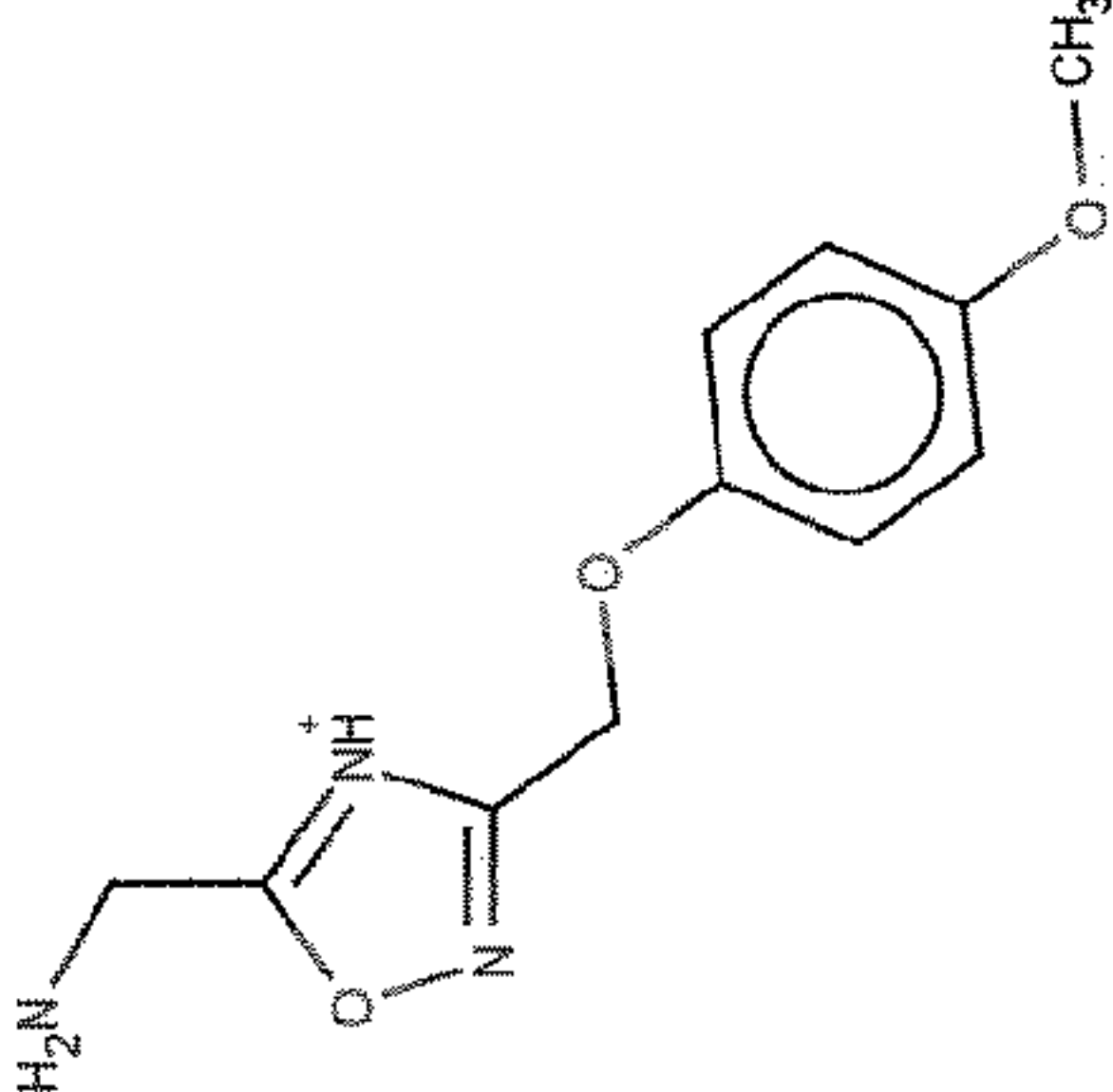
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 109		F2150-0040		F2150-0040_3_1 (1 of 4)
Compound 110		F2158-0133		F2158-0133_1_1 (1 of 7)
Compound 111		F2158-0625		F2158-0625_1 (1 of 7)

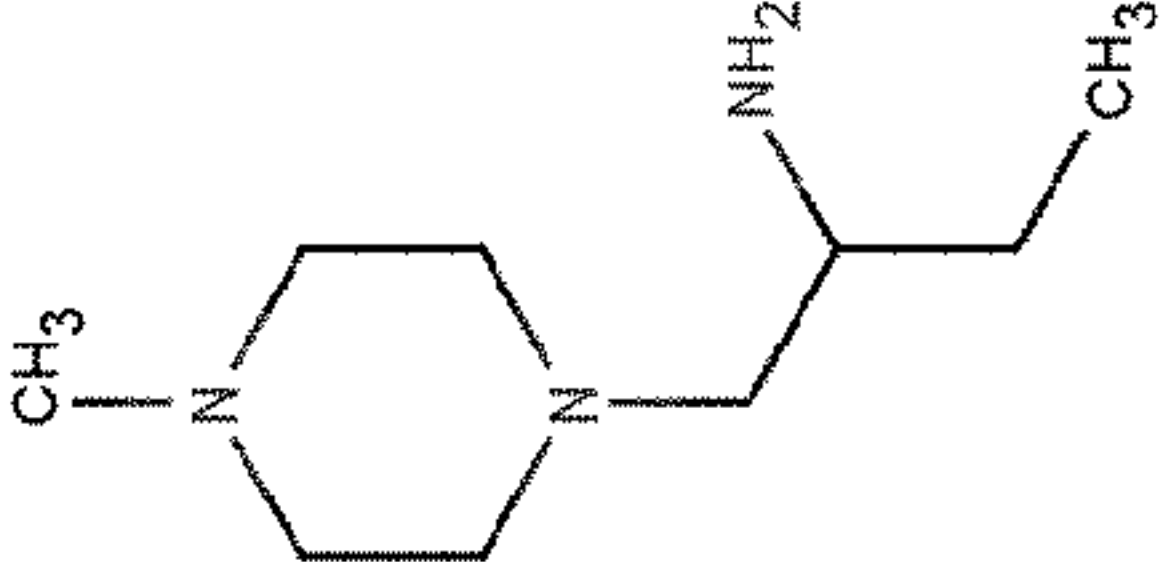
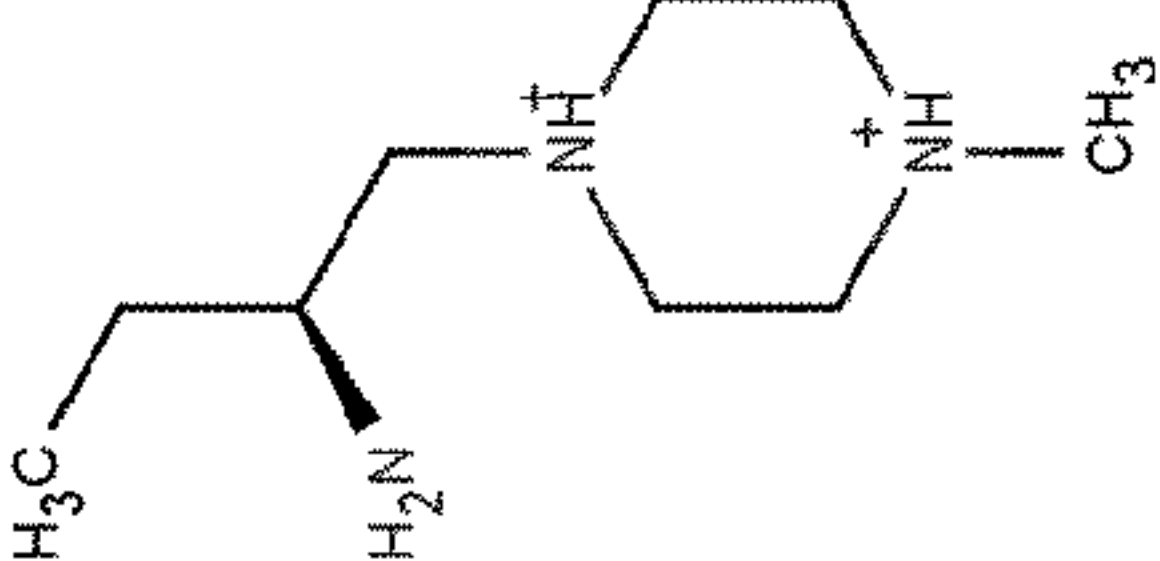
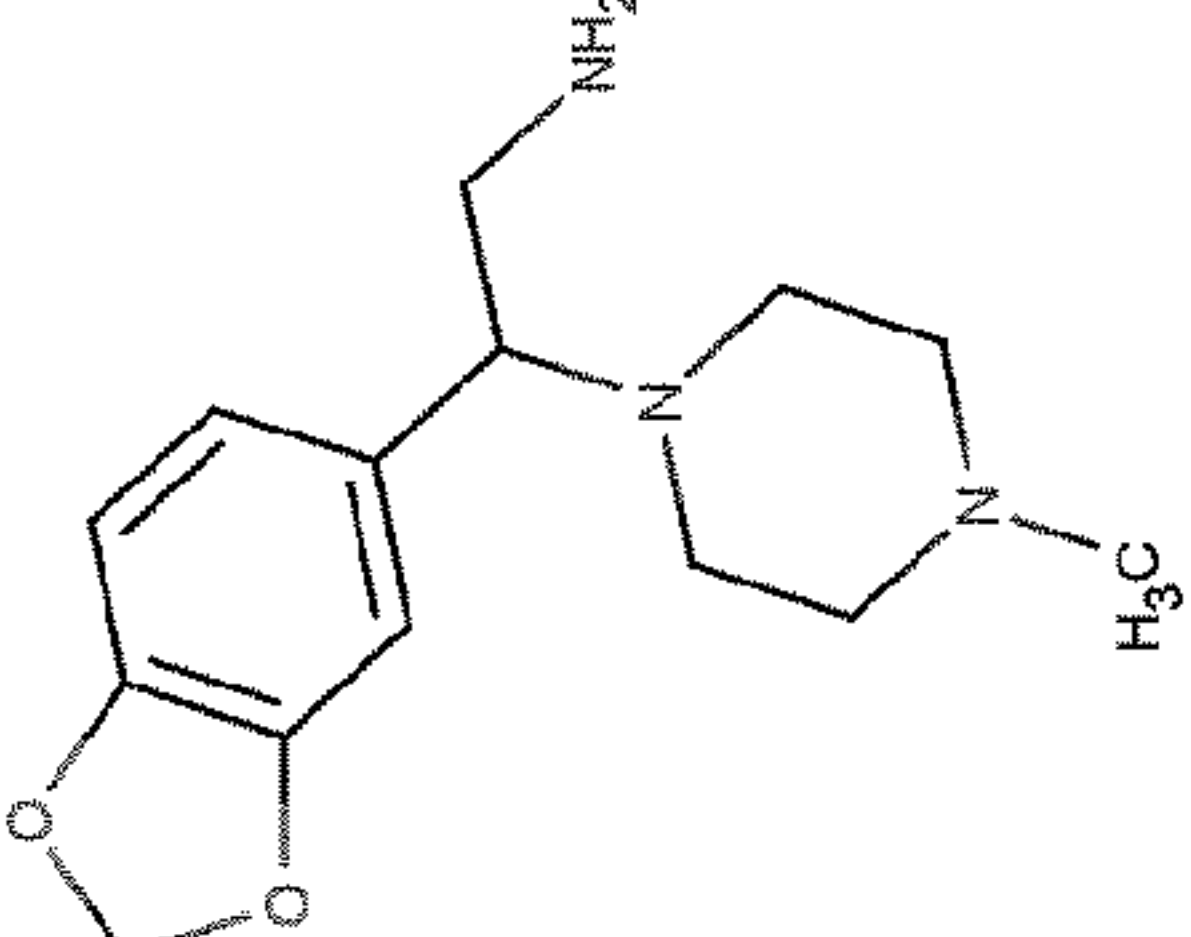
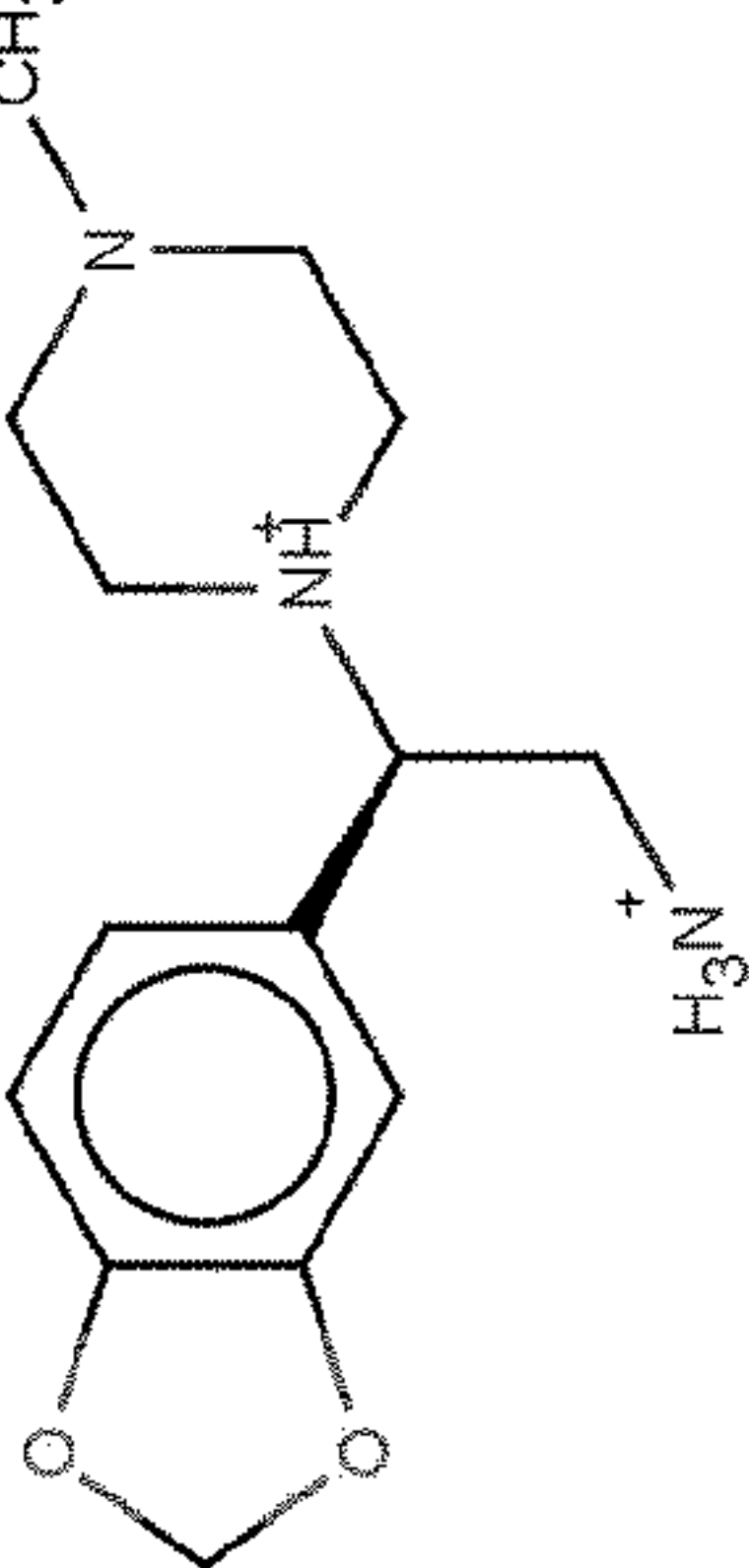
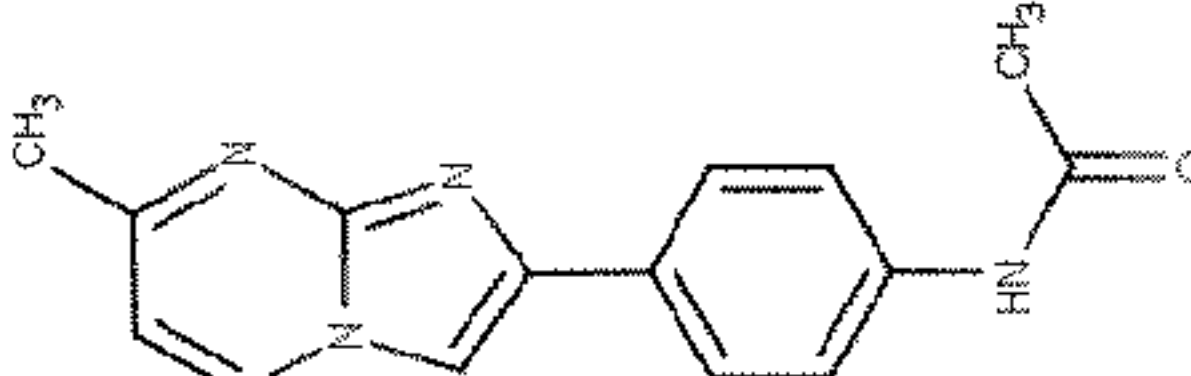
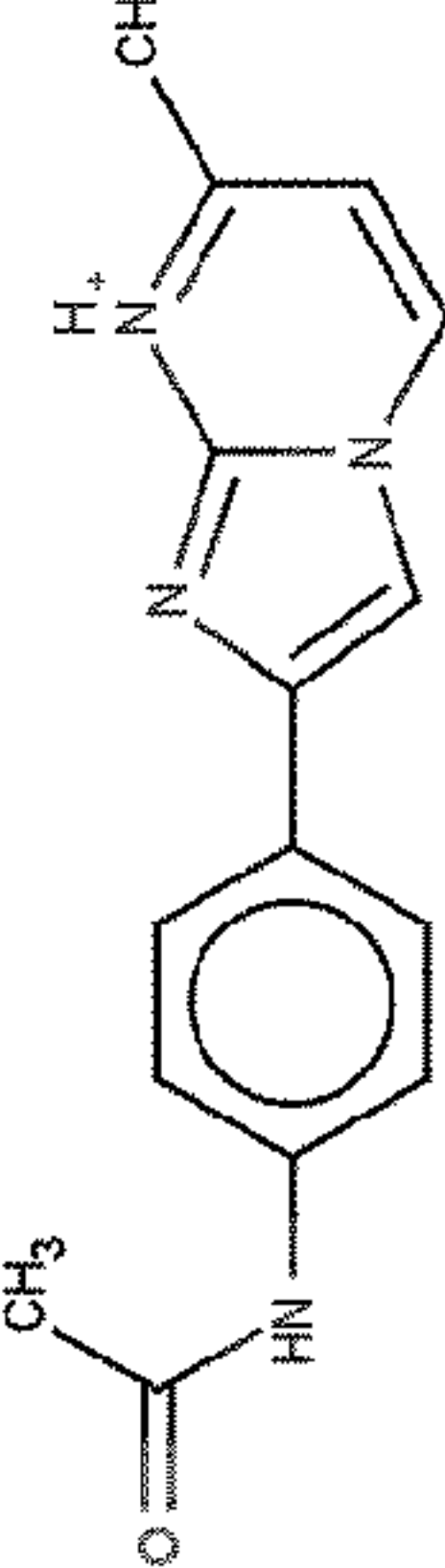
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 112		F2158-0833		F2158-0833_3_1 (1 of 4)
Compound 113		F2158-1073		F2158-1073_4_1 (1 of 7)
Compound 114		F2164-0018		F2164-0018_5_1 (1 of 7)

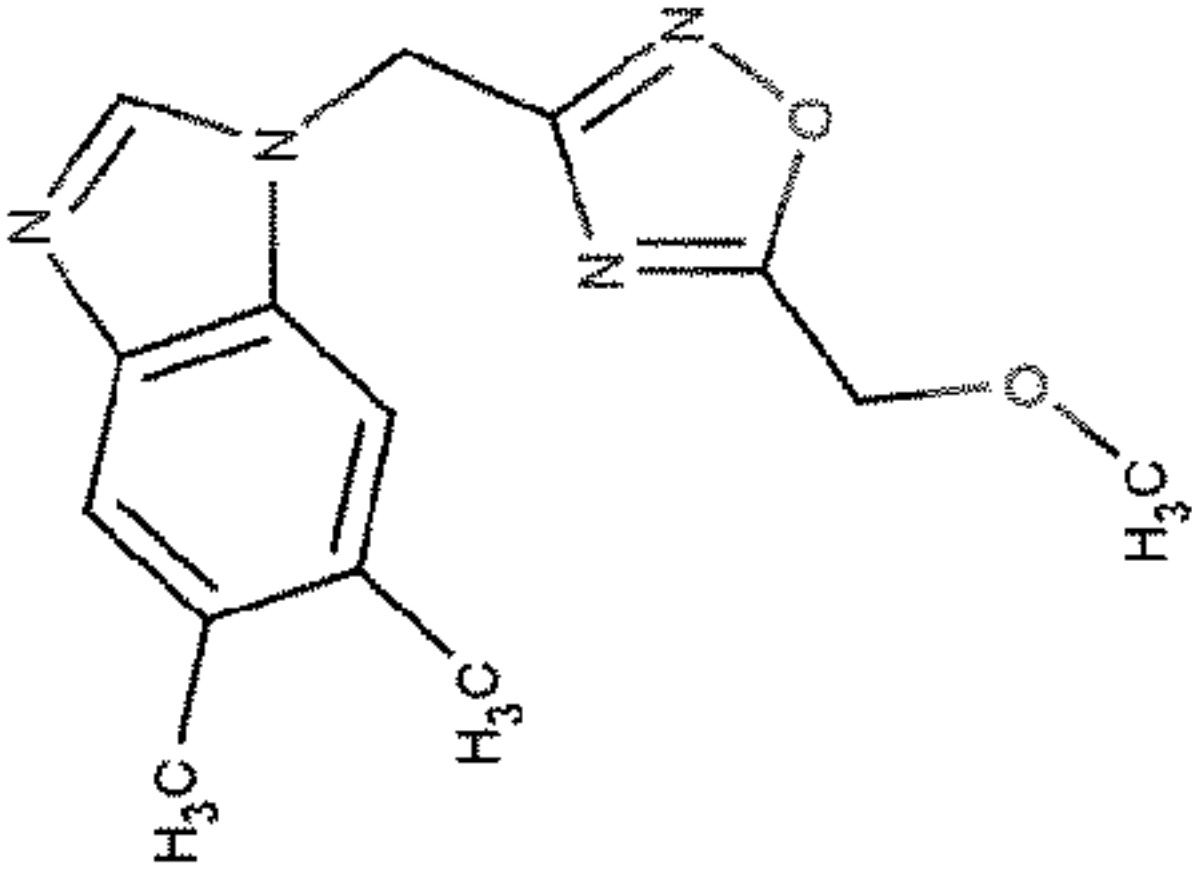
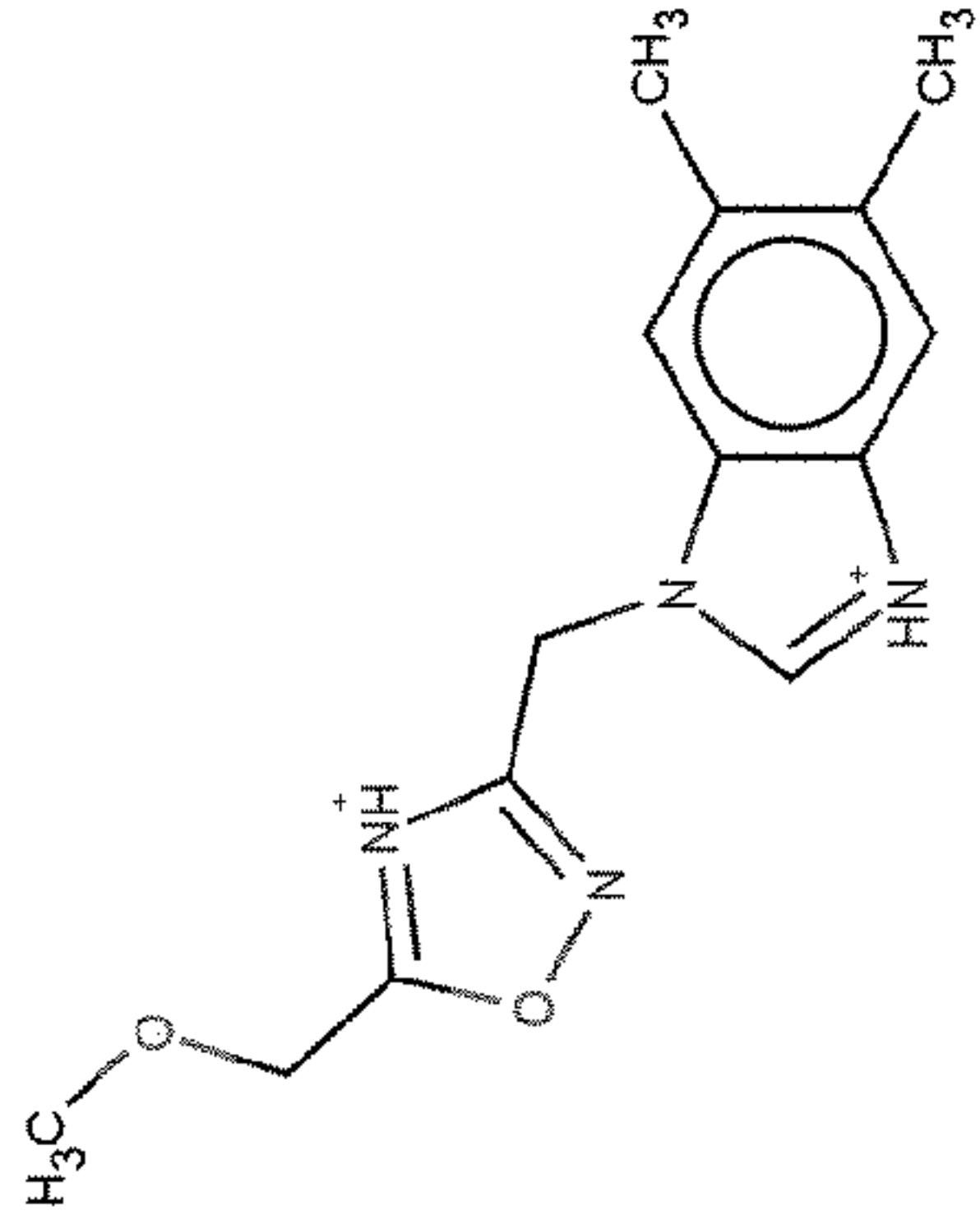
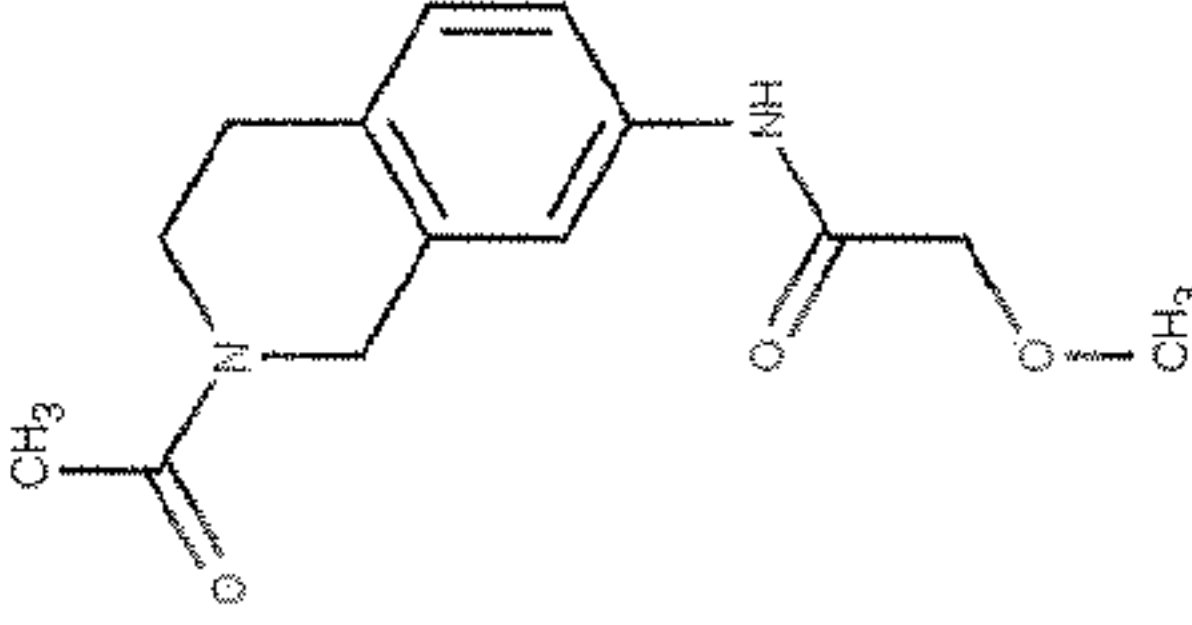
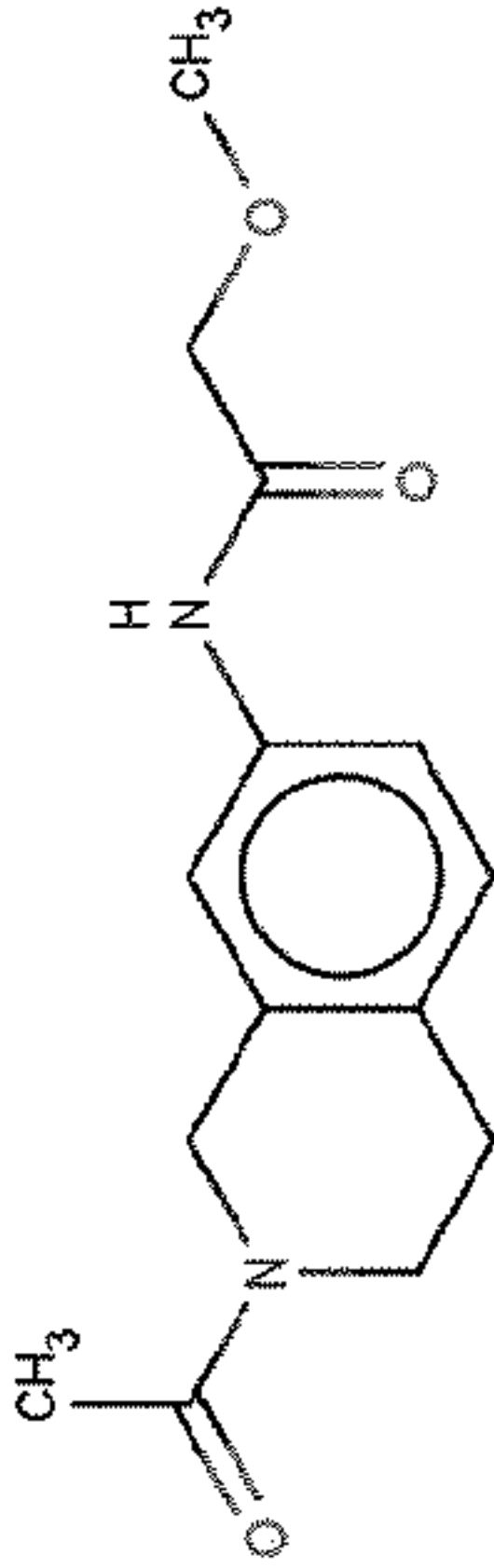
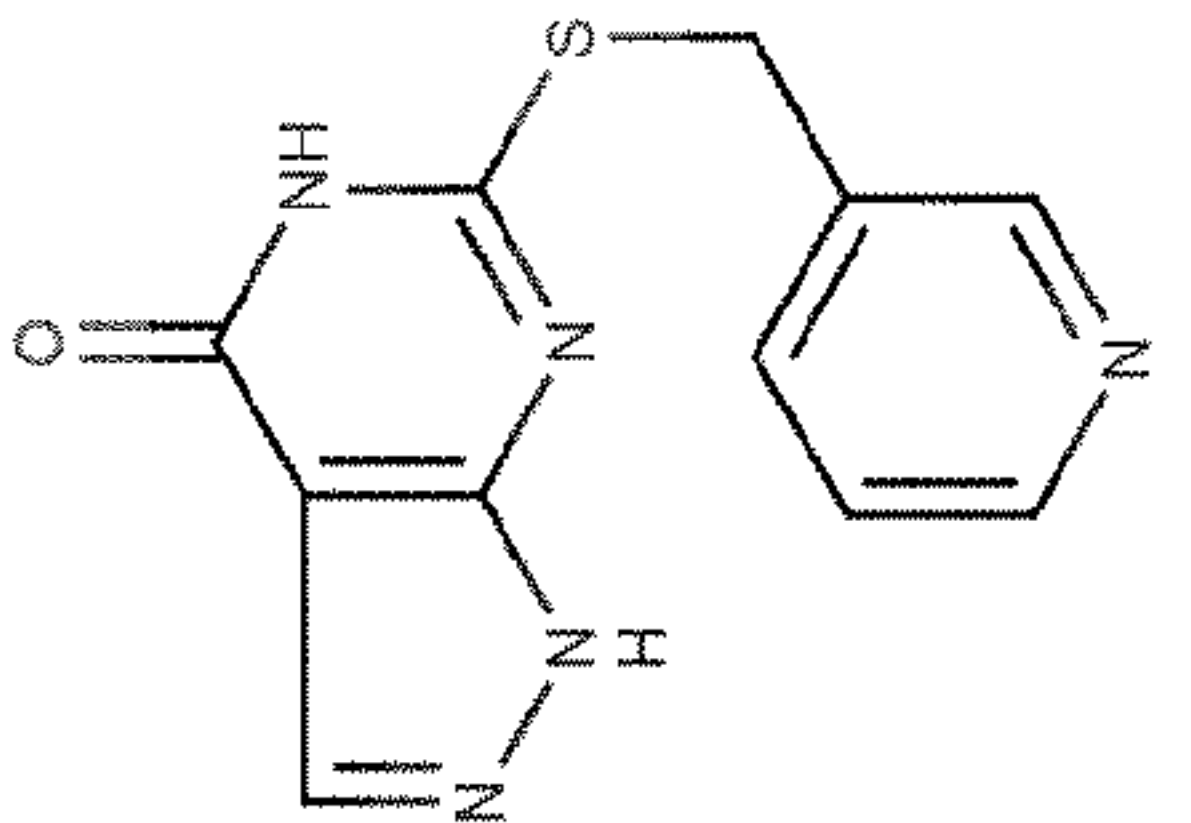
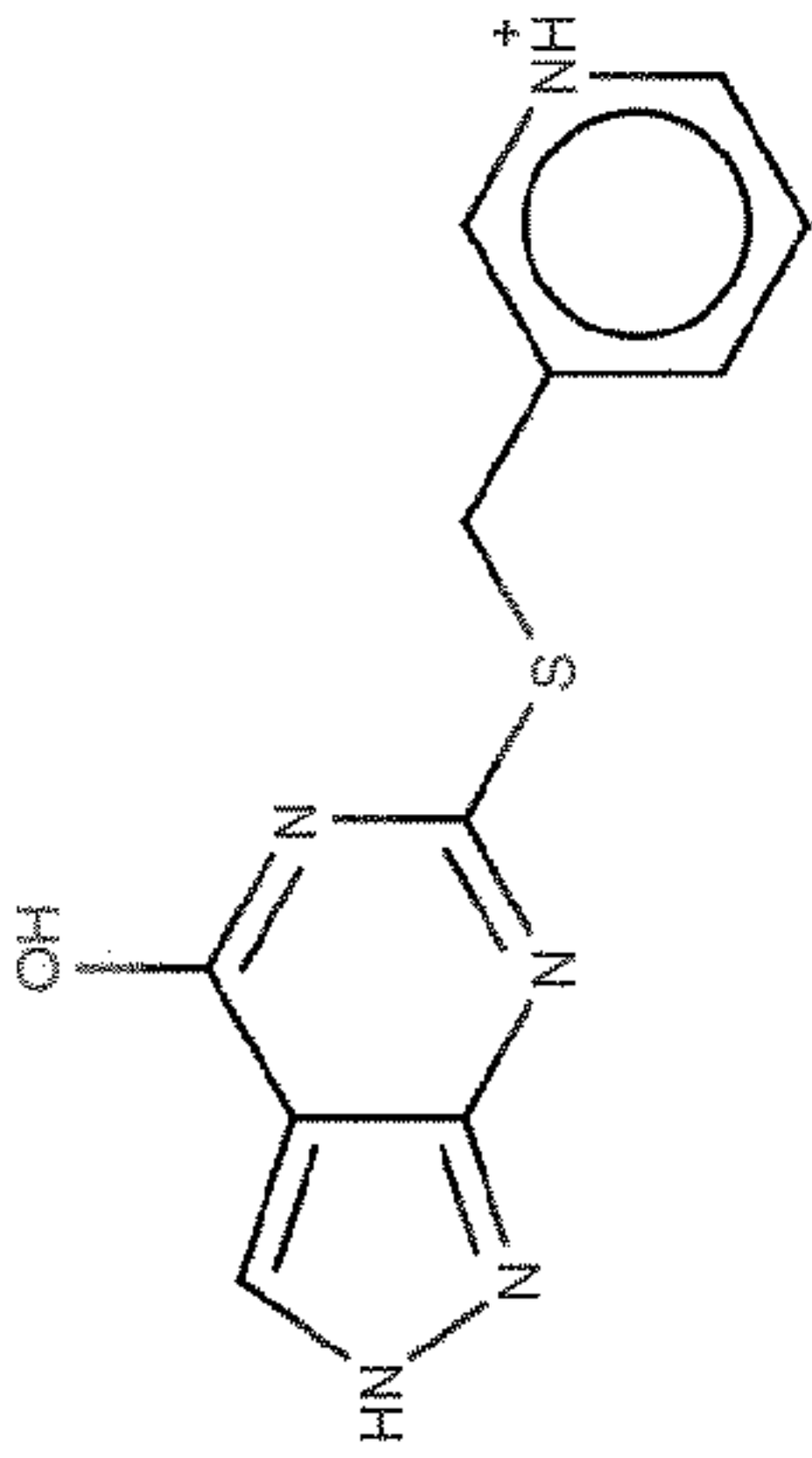
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 115		F2182-0033		F2182-0033_1_2 (1 of 4)
Compound 116		F2182-0073		F2182-0073_3_1 (1 of 4)
Compound 117		F2182-0075		F2182-0075_3_1 (1 of 4)

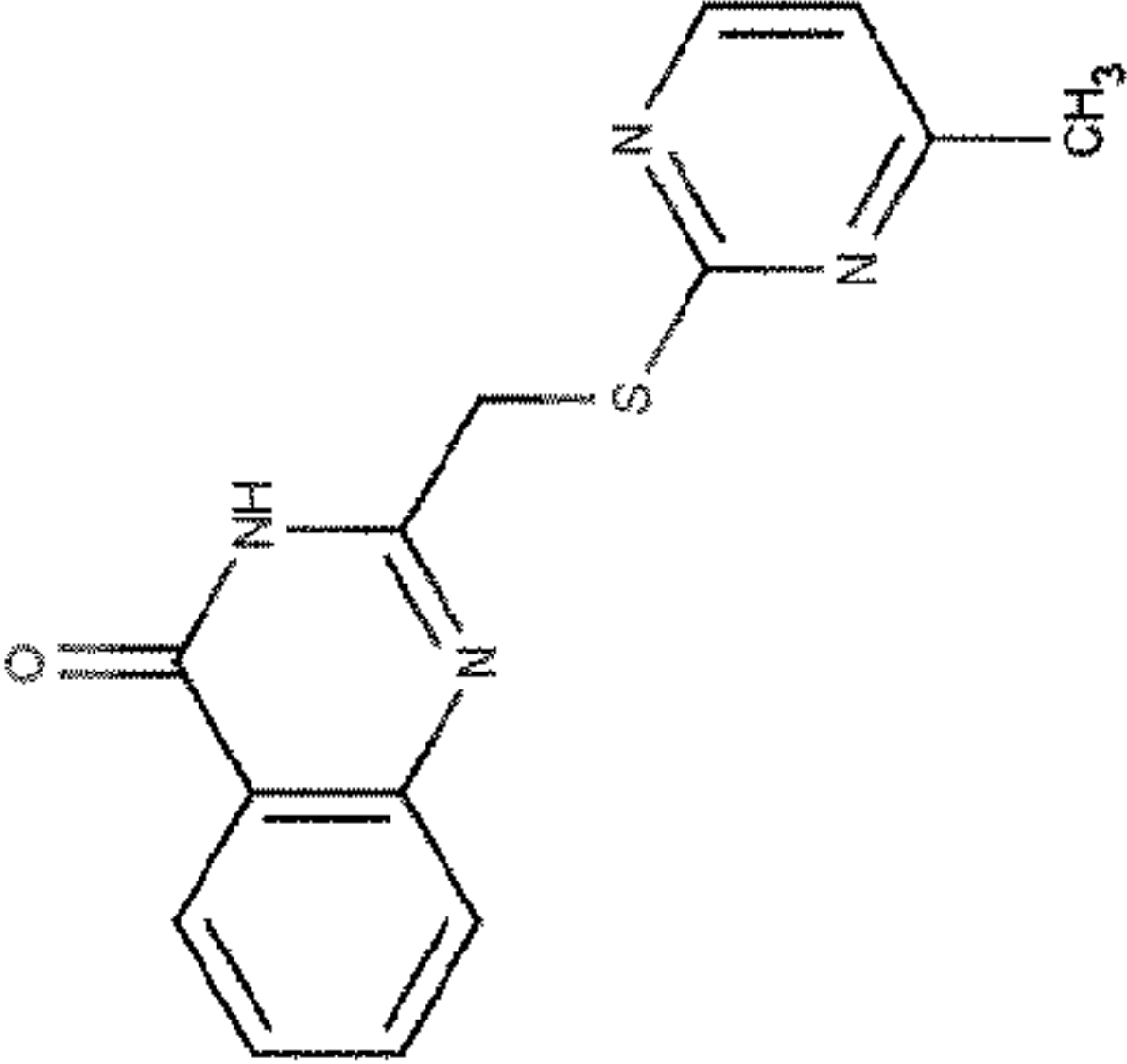
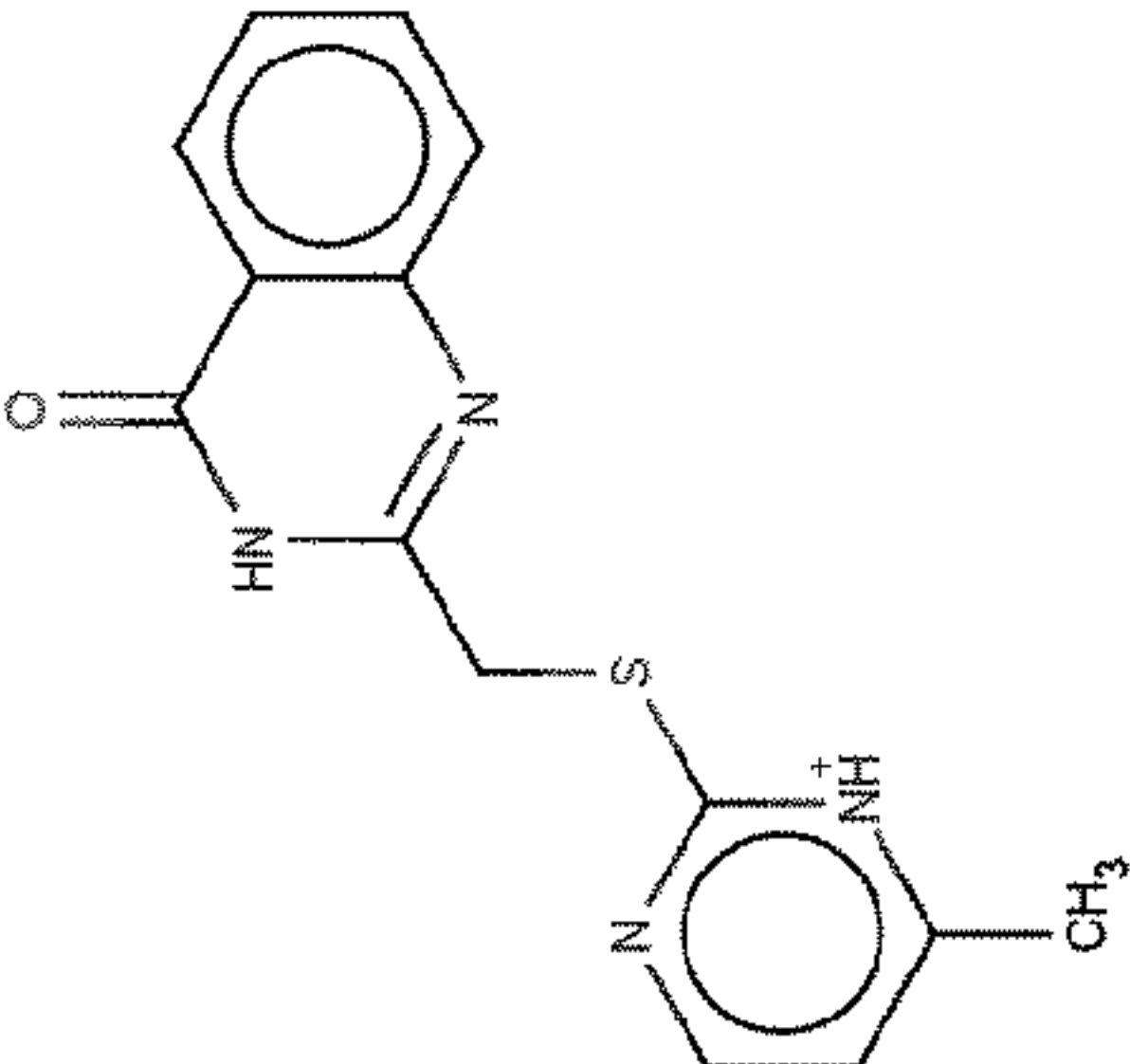
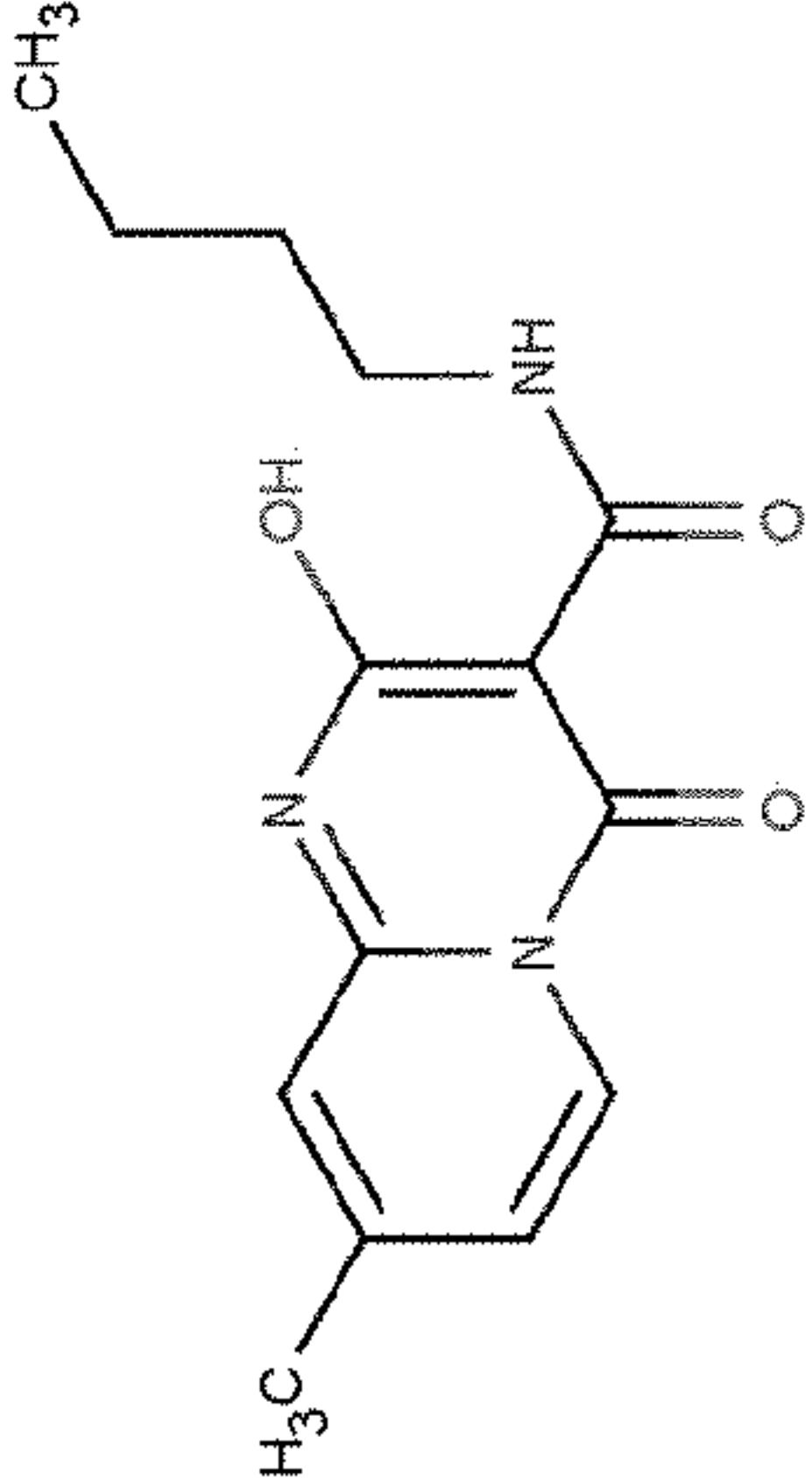
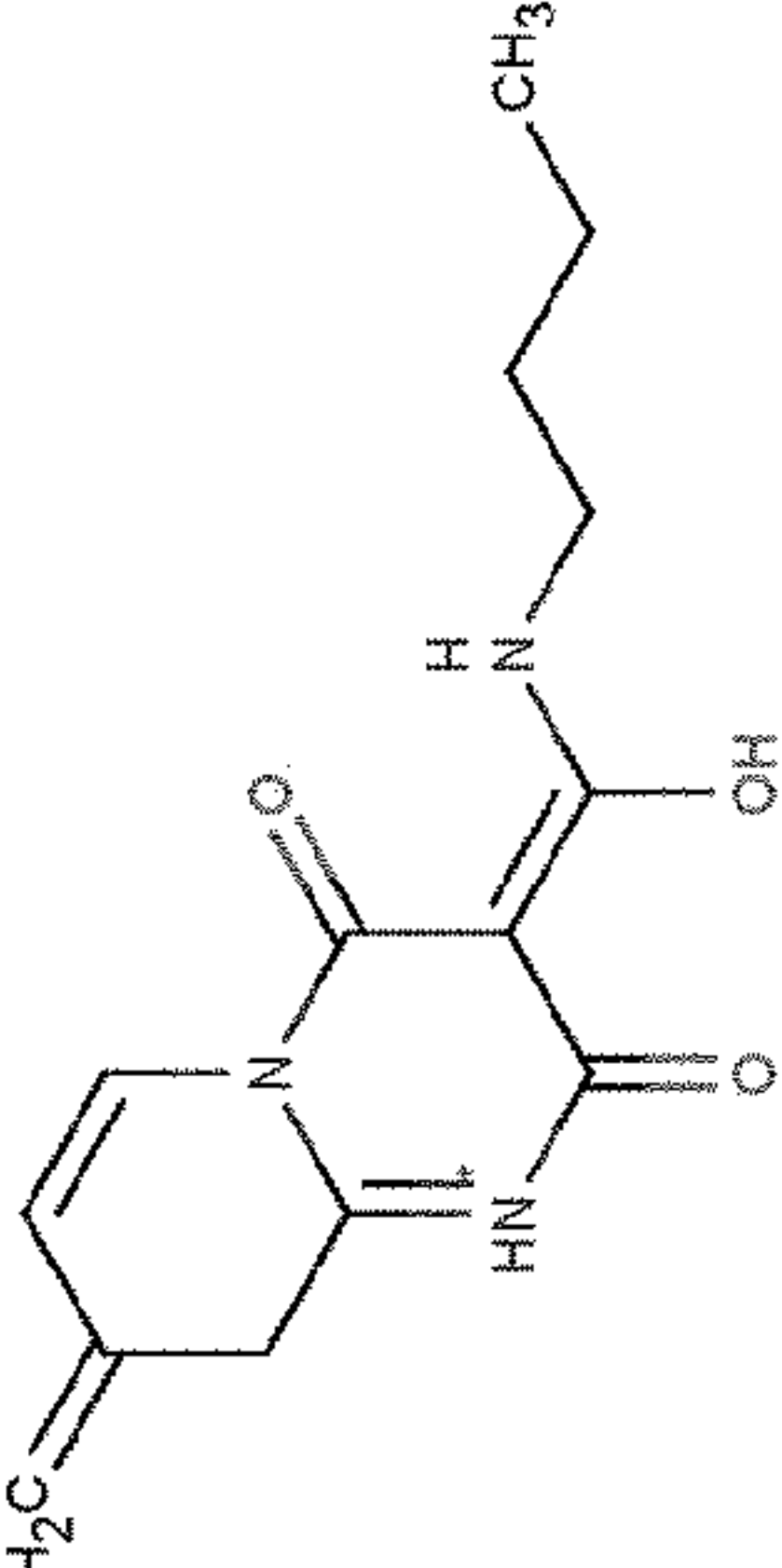
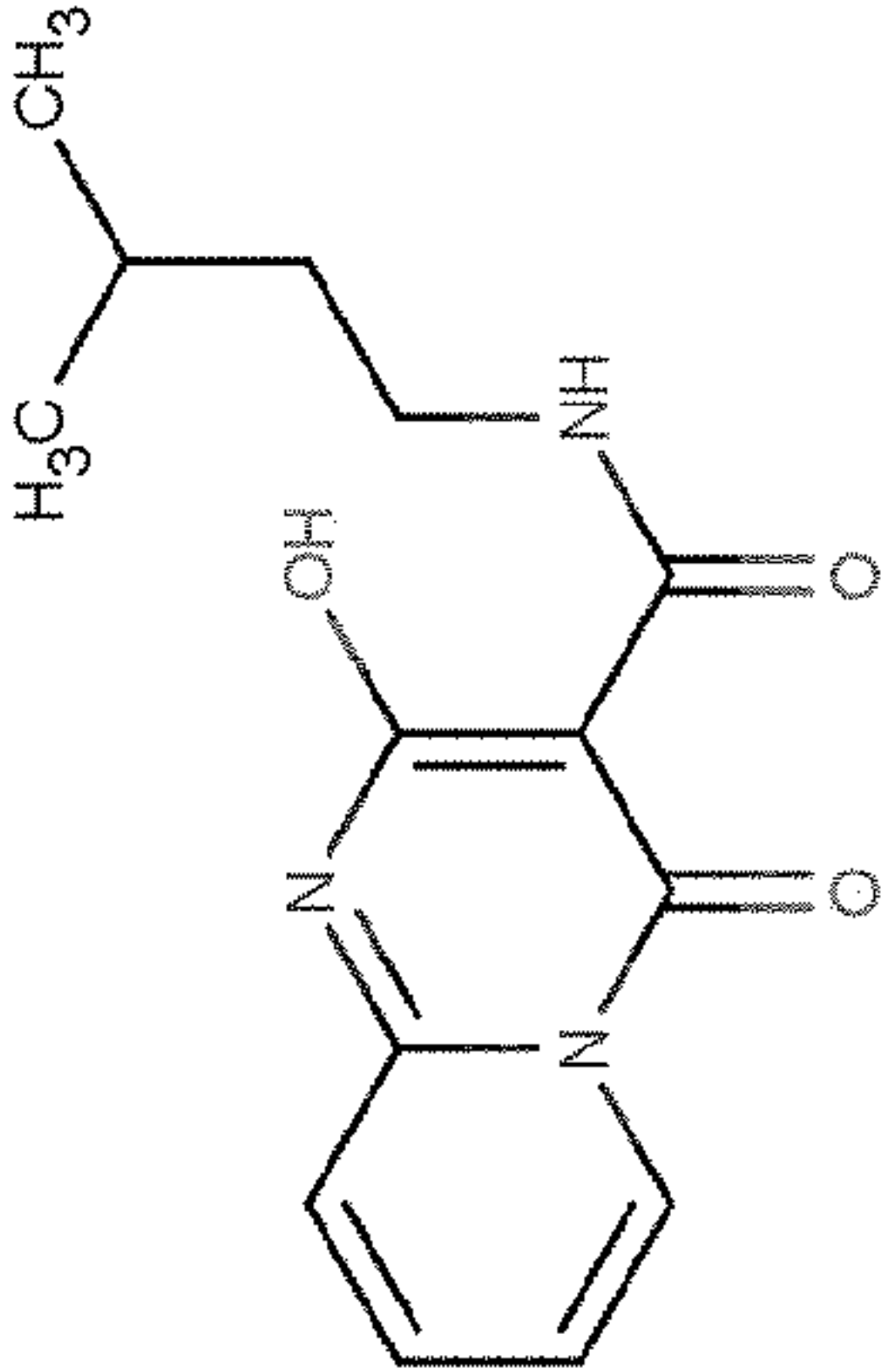
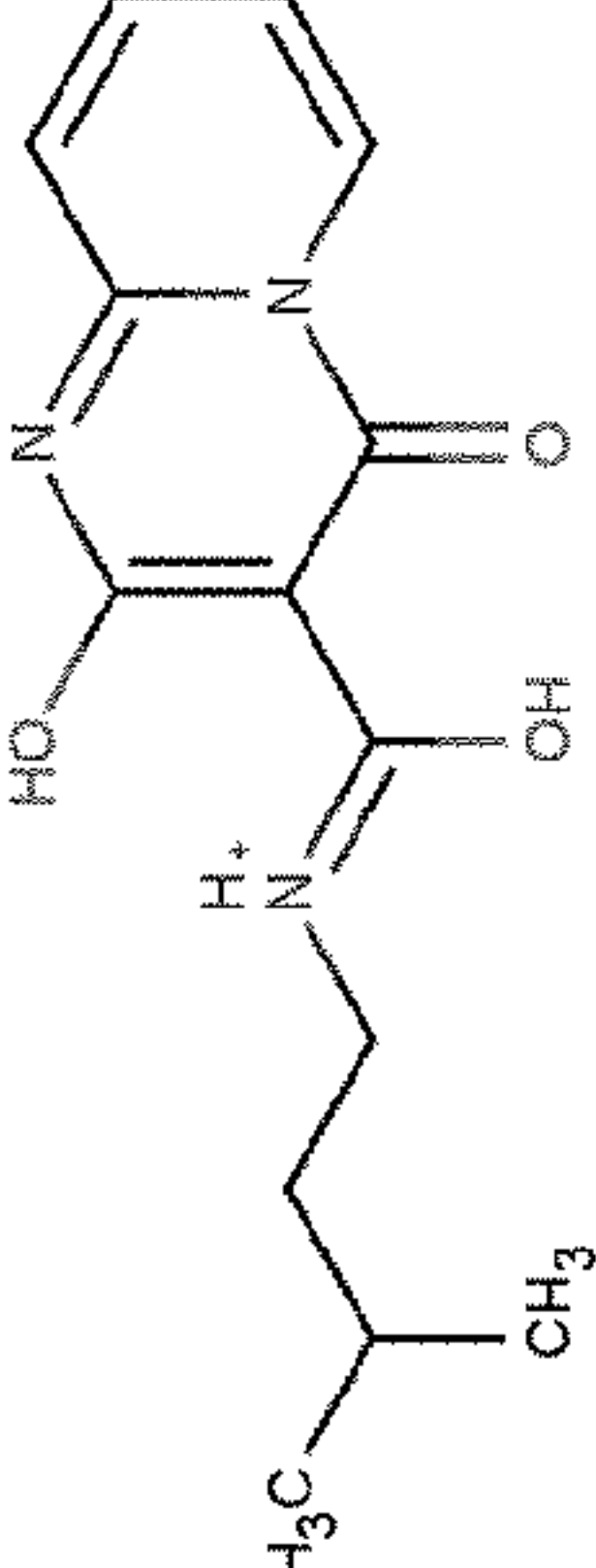
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 118		F2182-0085		F2182-0085_1_1 (1 of 4)
Compound 119		F2182-0091		F2182-0091_1_1 (1 of 4)
Compound 120		F2182-0097		F2182-0097_1_1 (1 of 4)

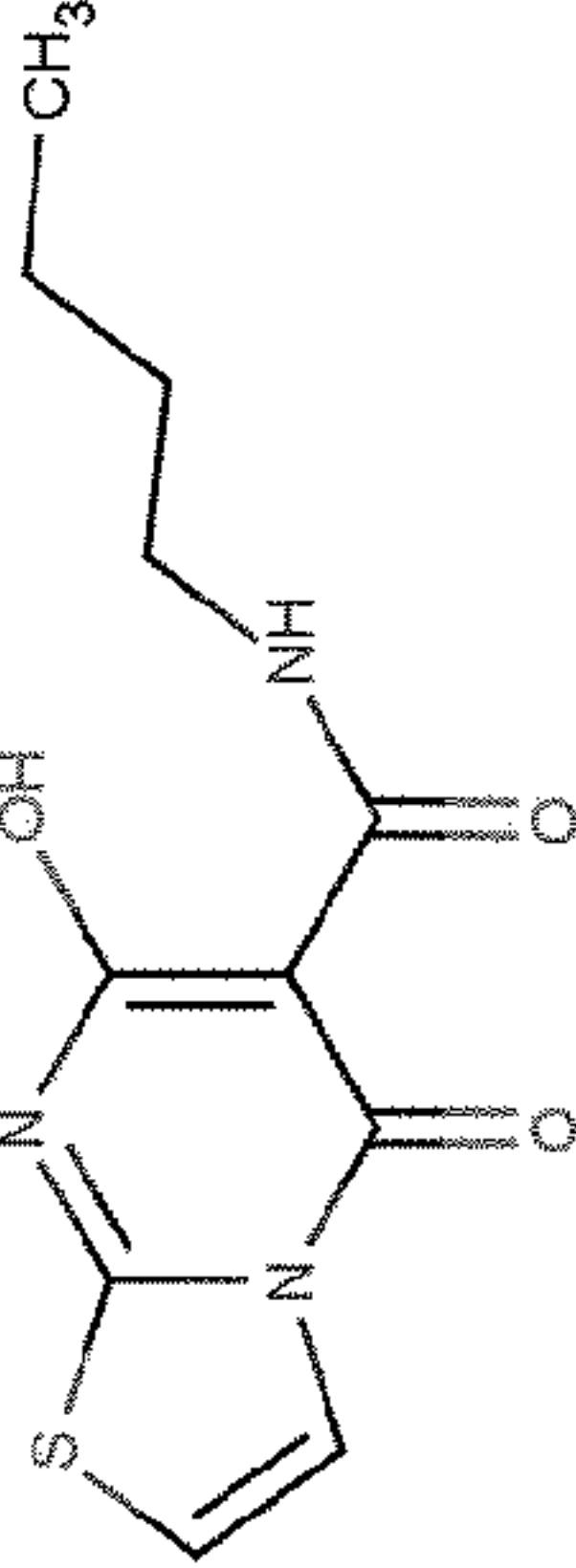
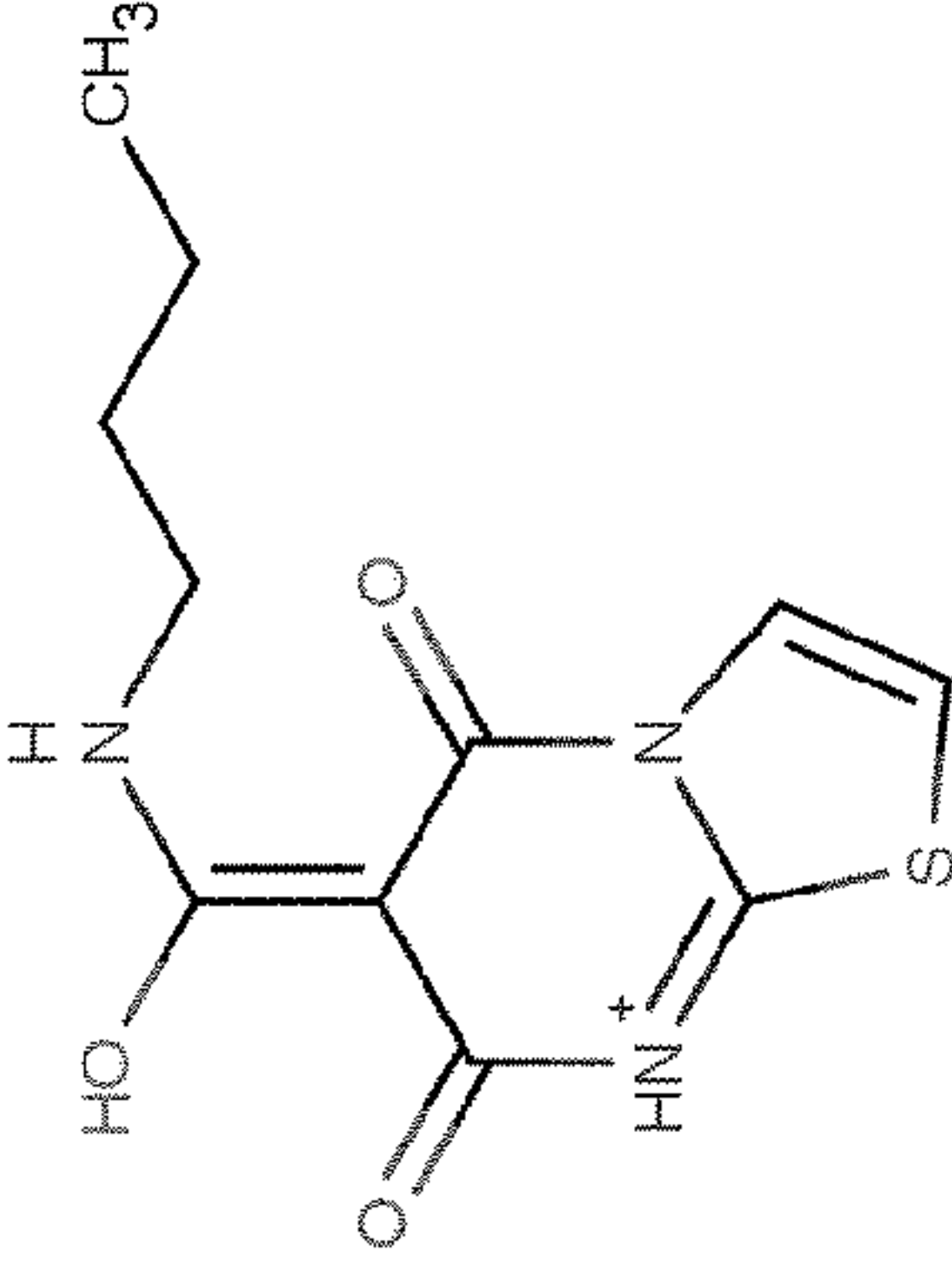
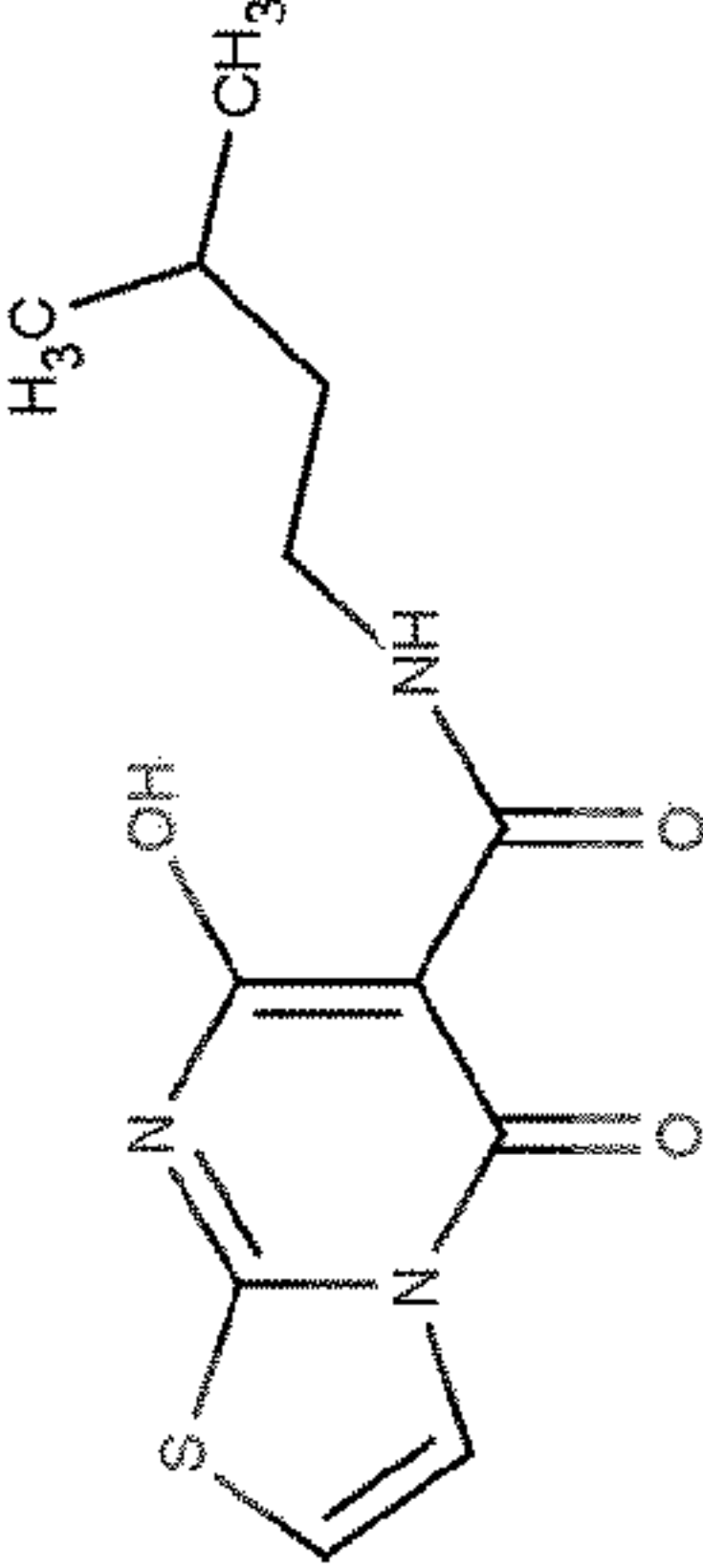
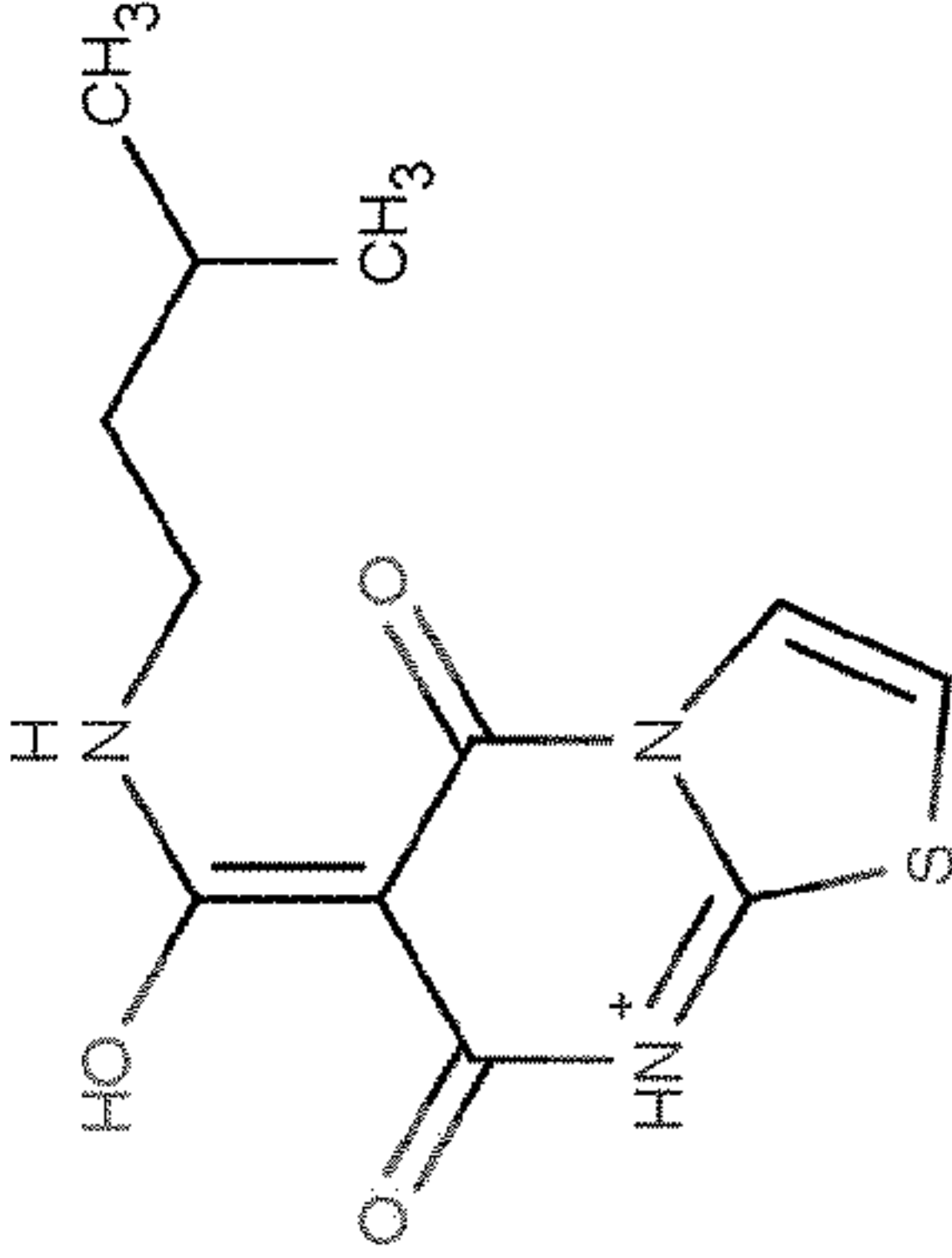
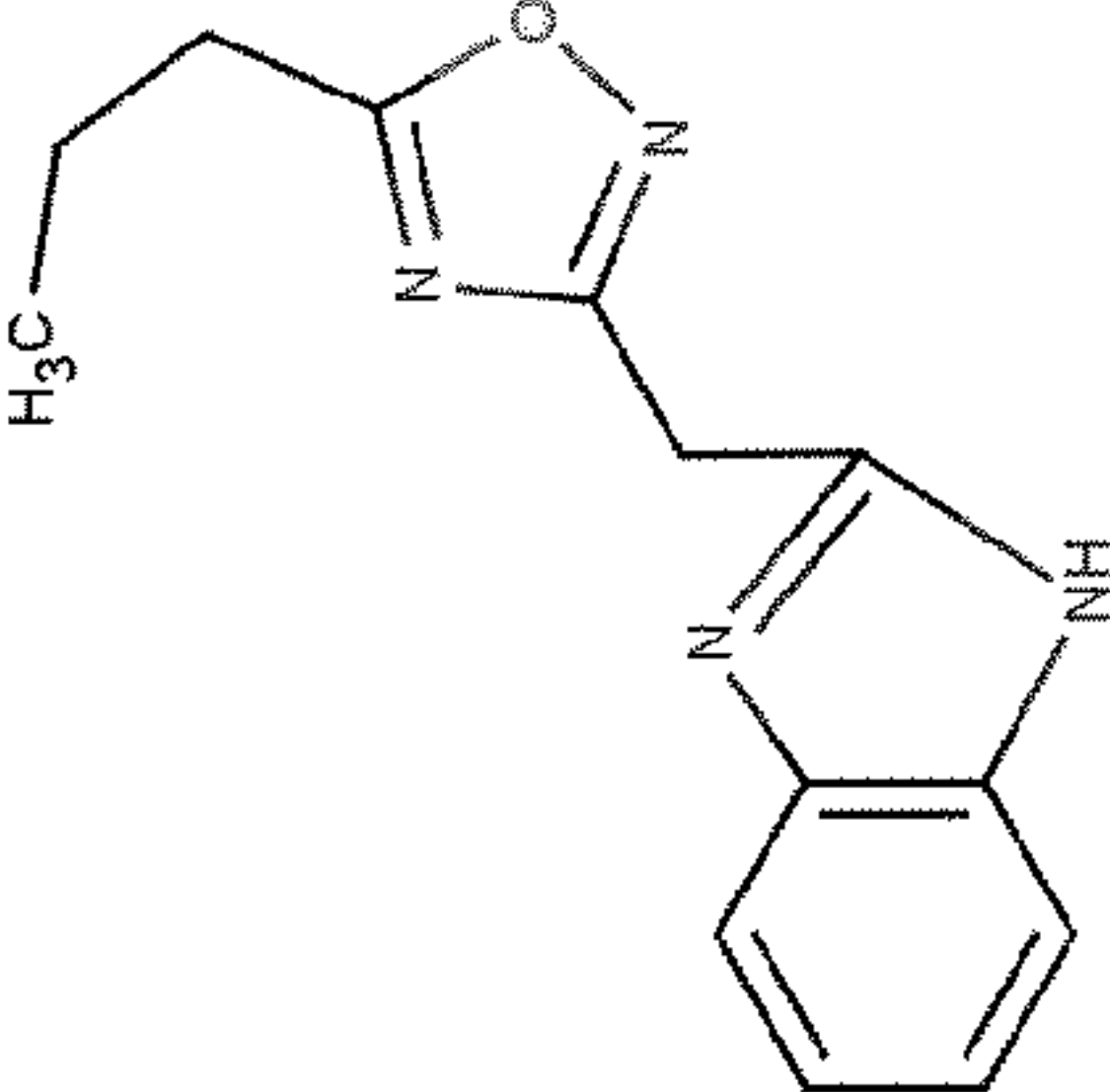
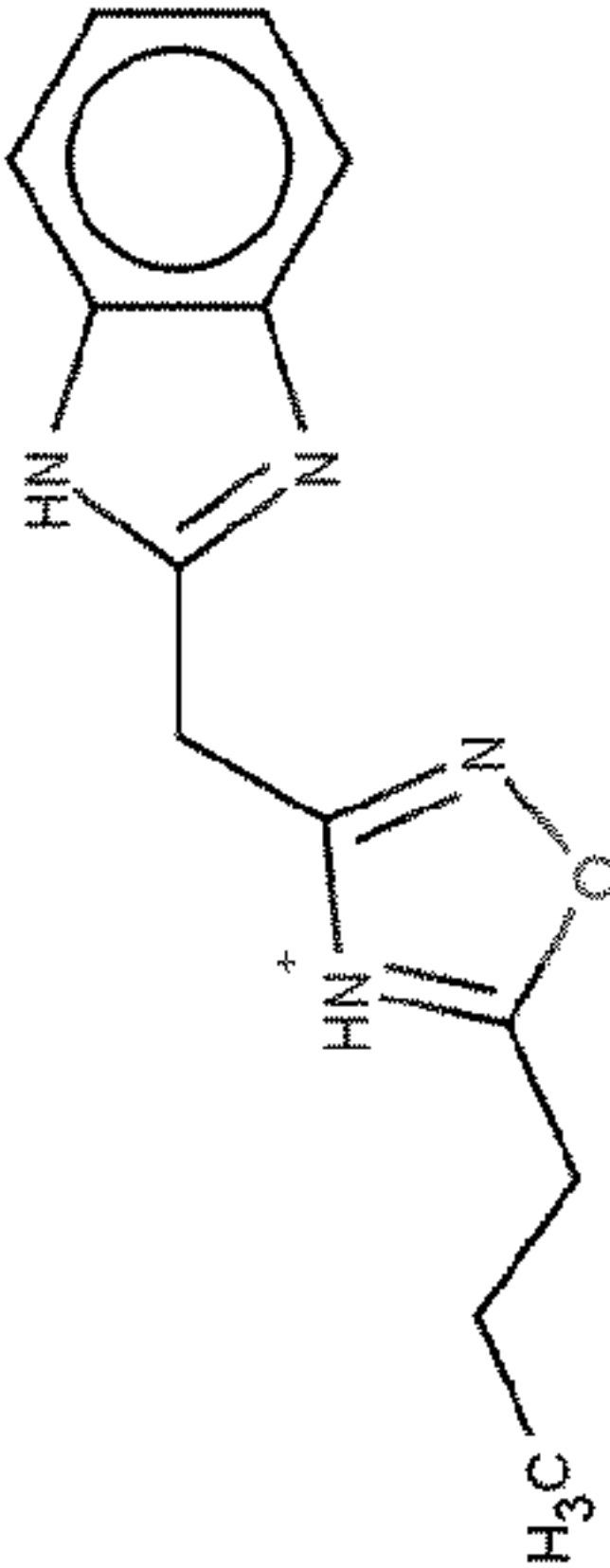
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 121		F2182-0103		F2182-0103_3_1 (1 of 4)
Compound 122		F2182-0106		F2182-0106_3_1 (1 of 4)
Compound 123		F2182-0107		F2182-0107_1_1 (1 of 4)

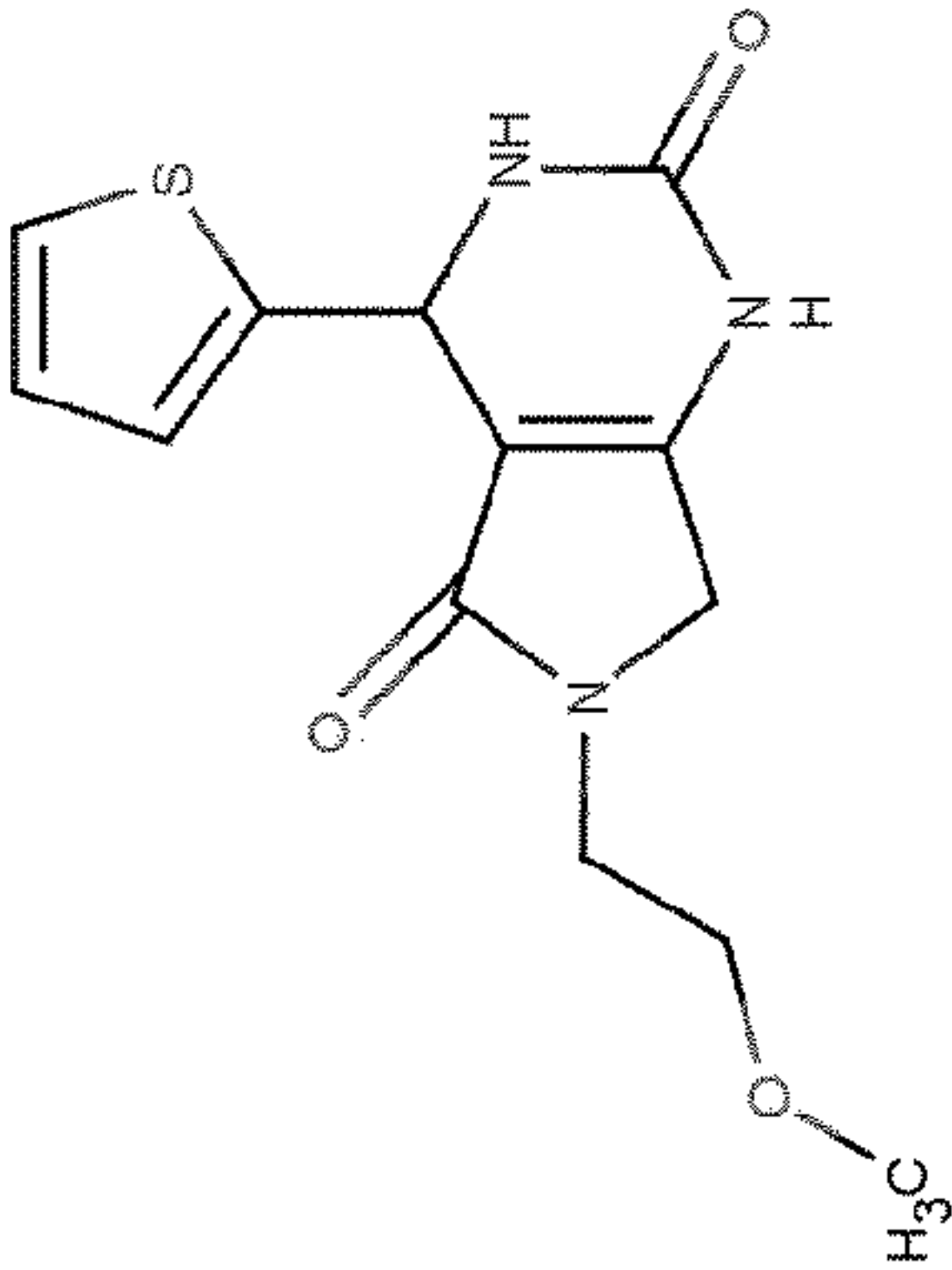
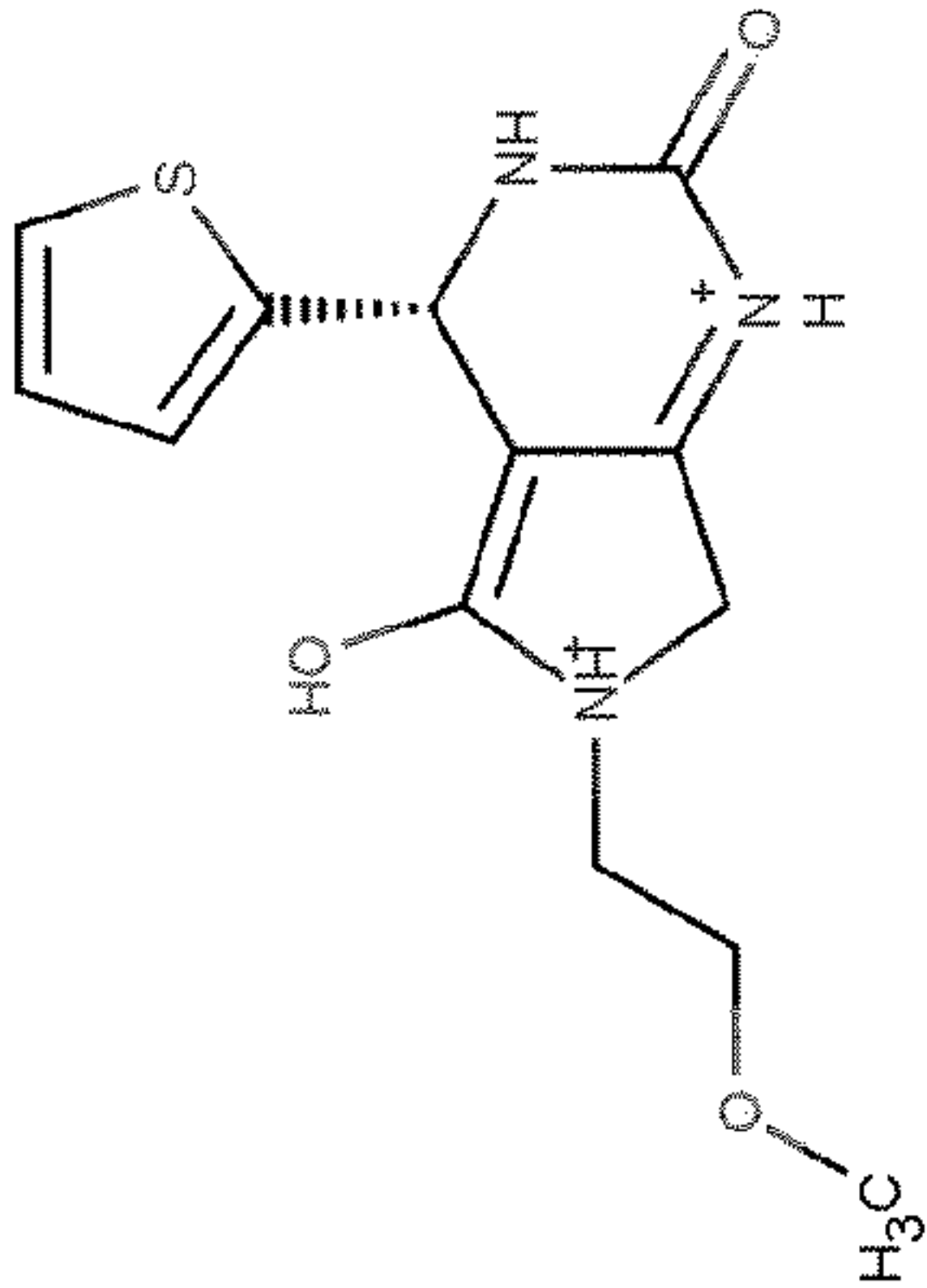
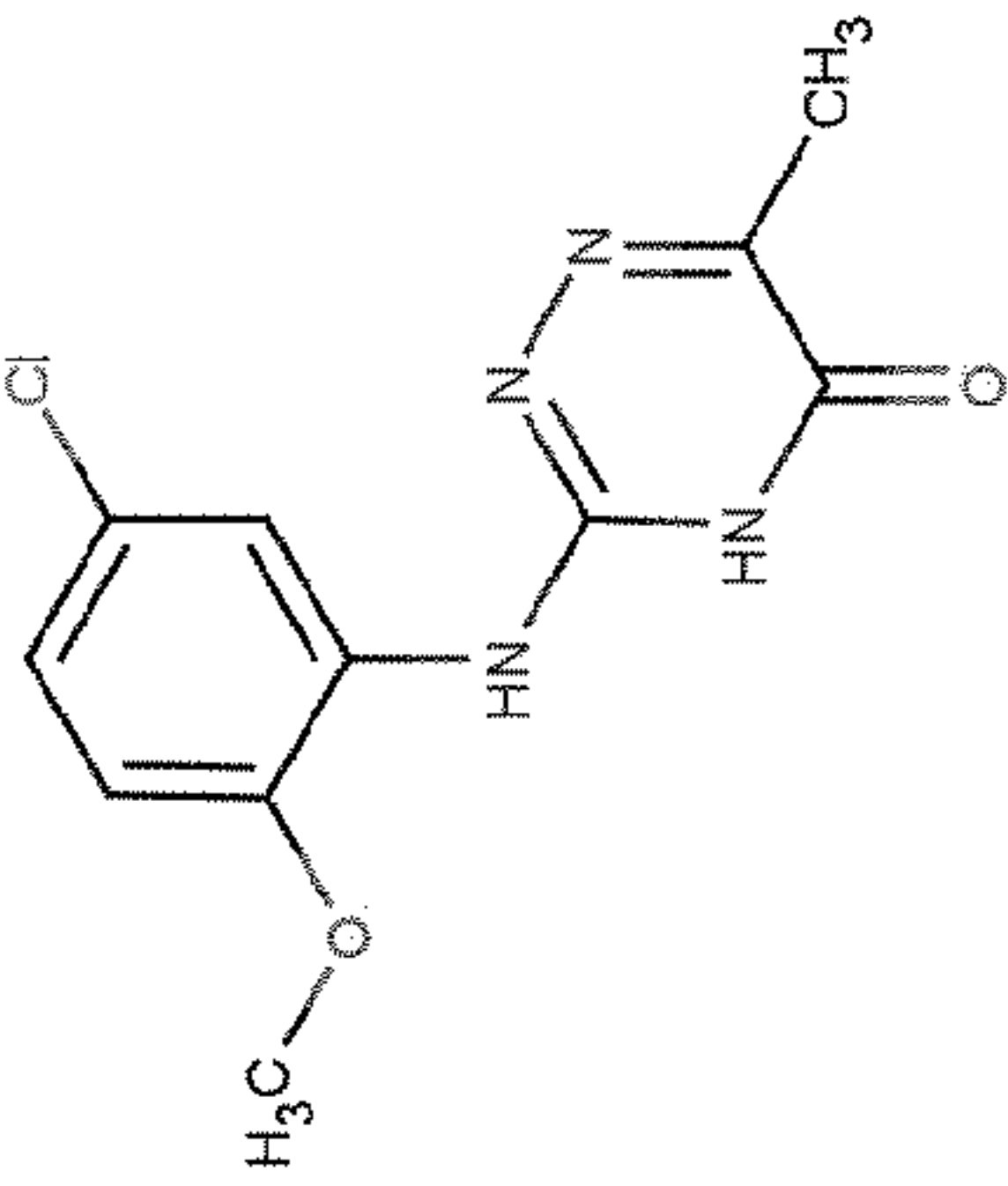
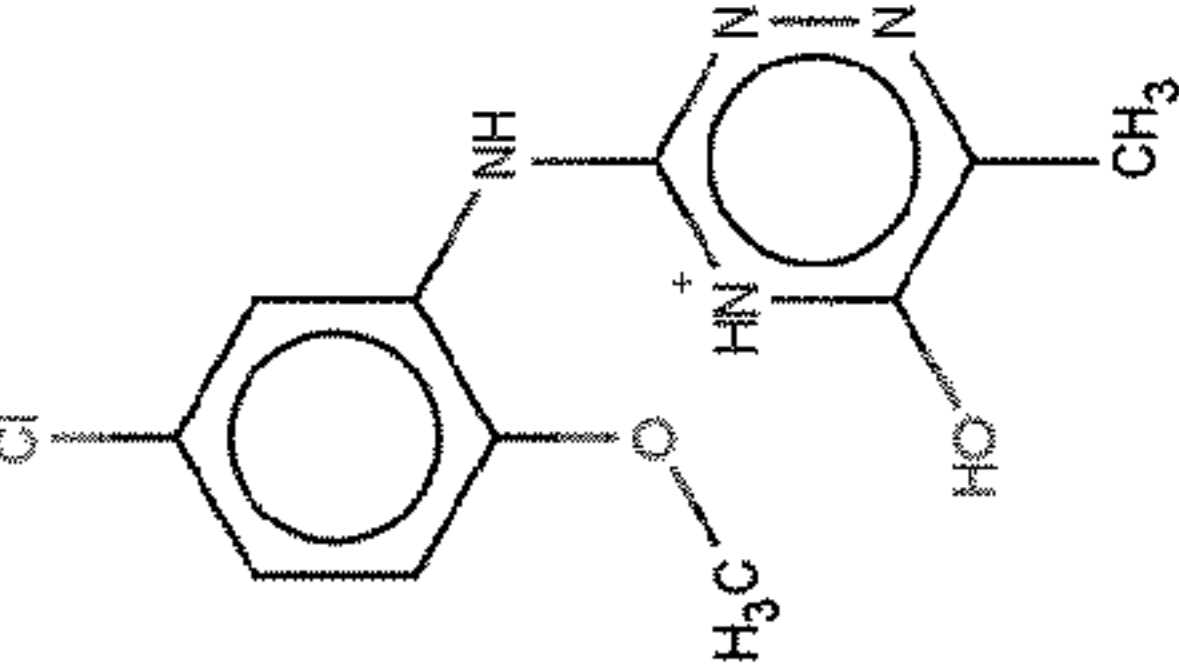
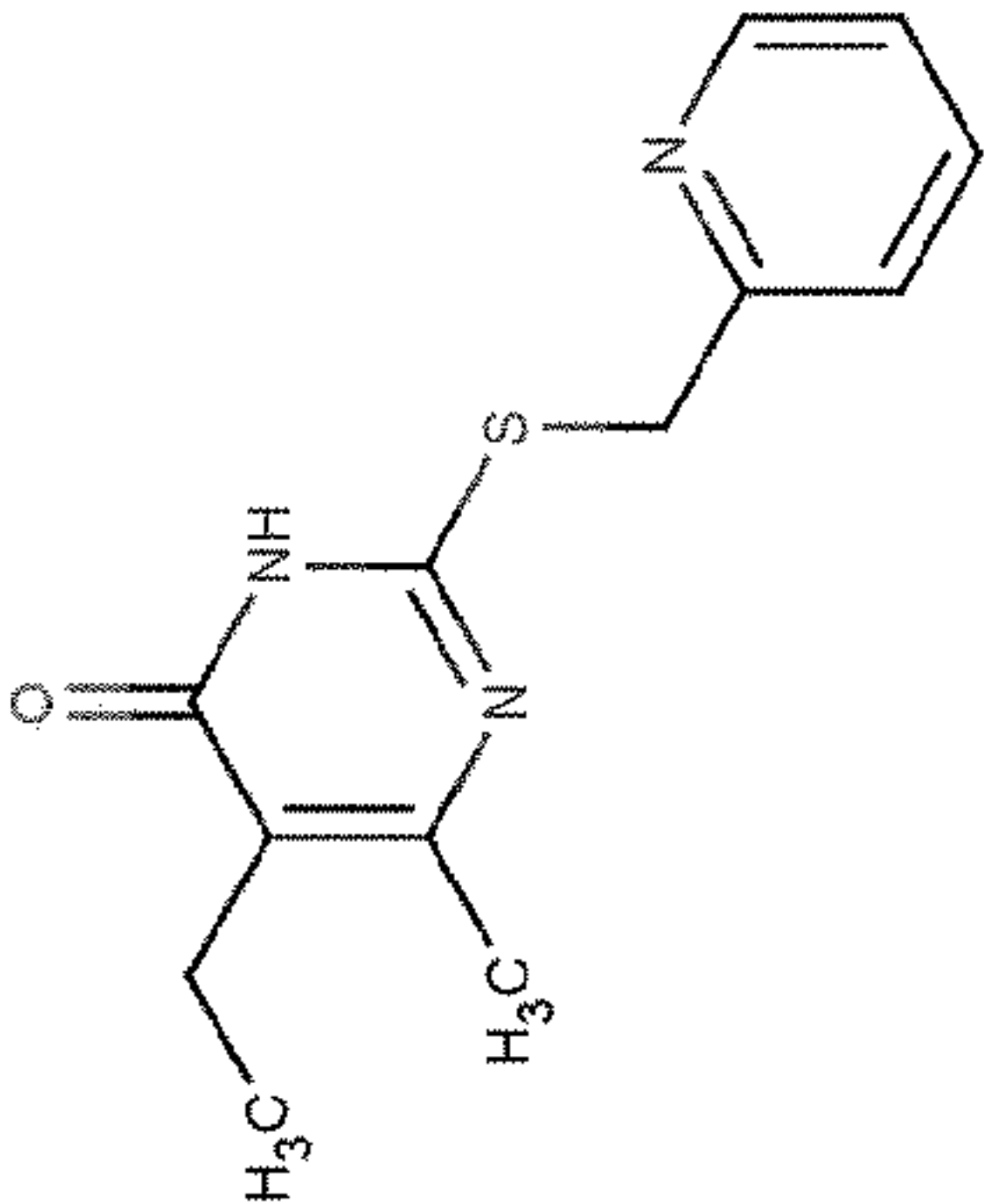
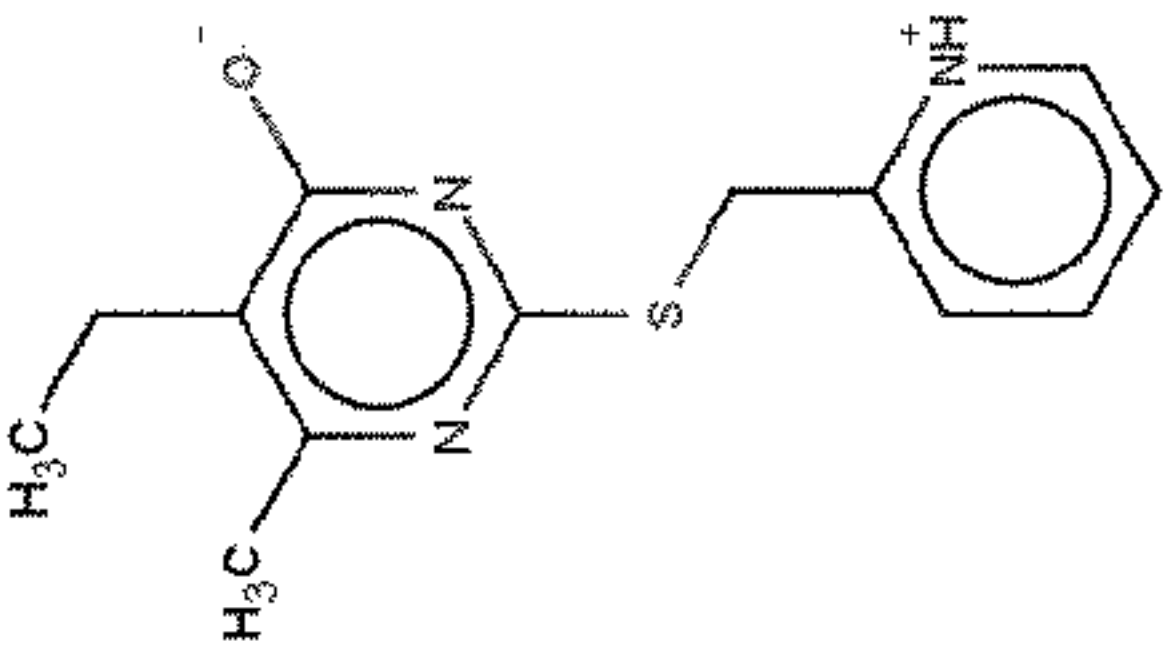
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 124		F2182-0112		F2182-0112_1_1 (1 of 4)
Compound 125		F2185-0004		F2185-0004_3_1 (1 of 4)
Compound 126		F2185-0006		F2185-0006_1_1 (1 of 4)

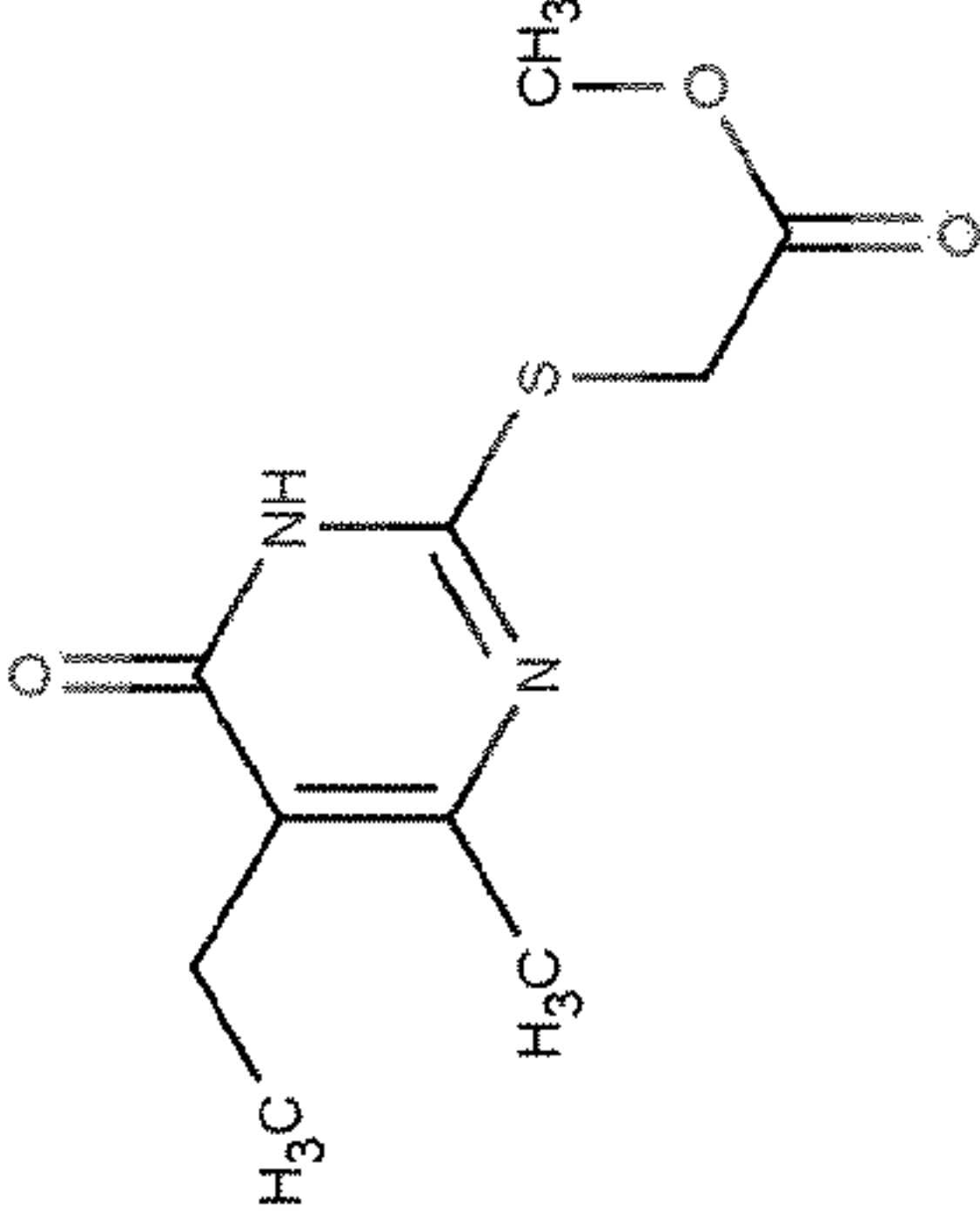
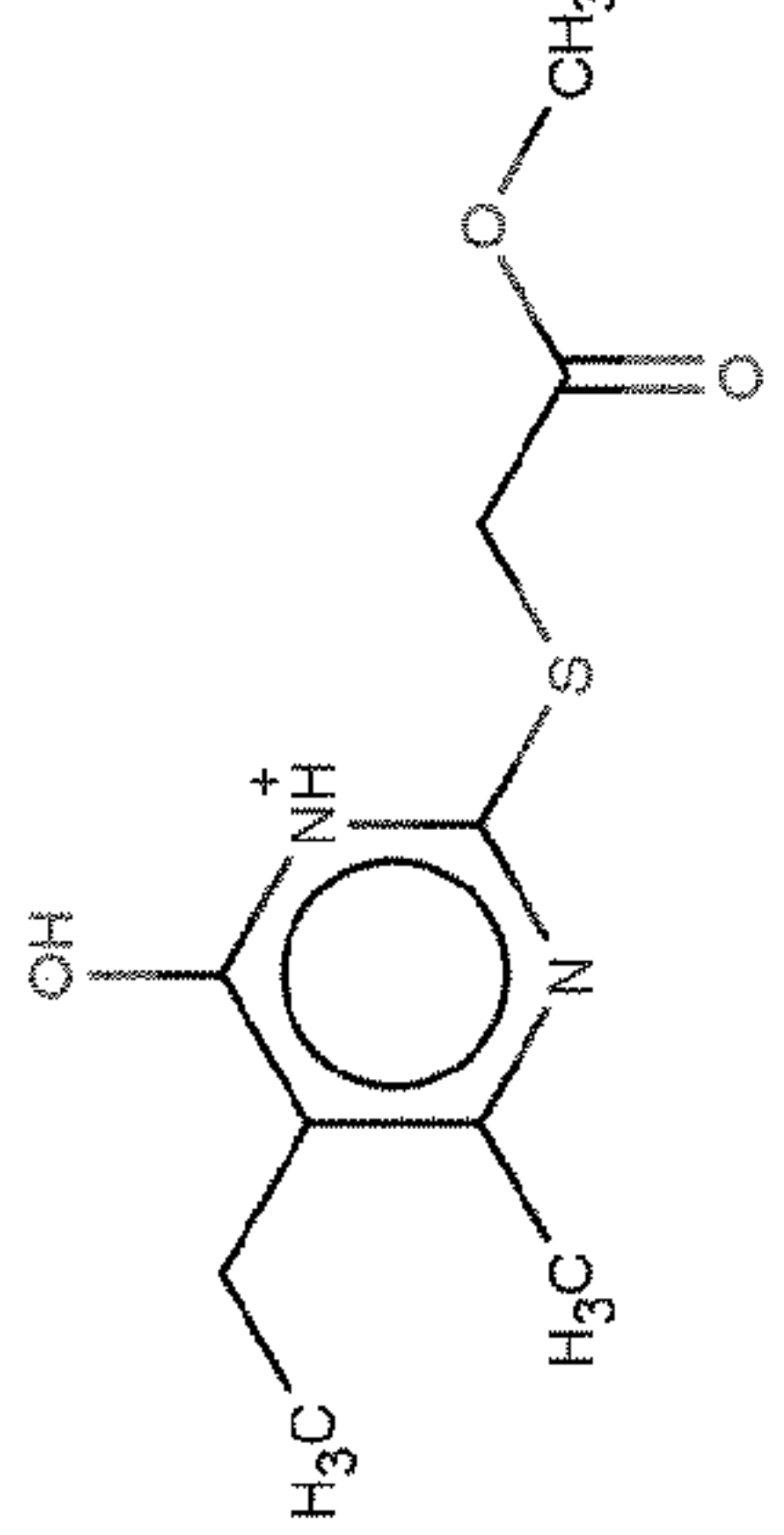
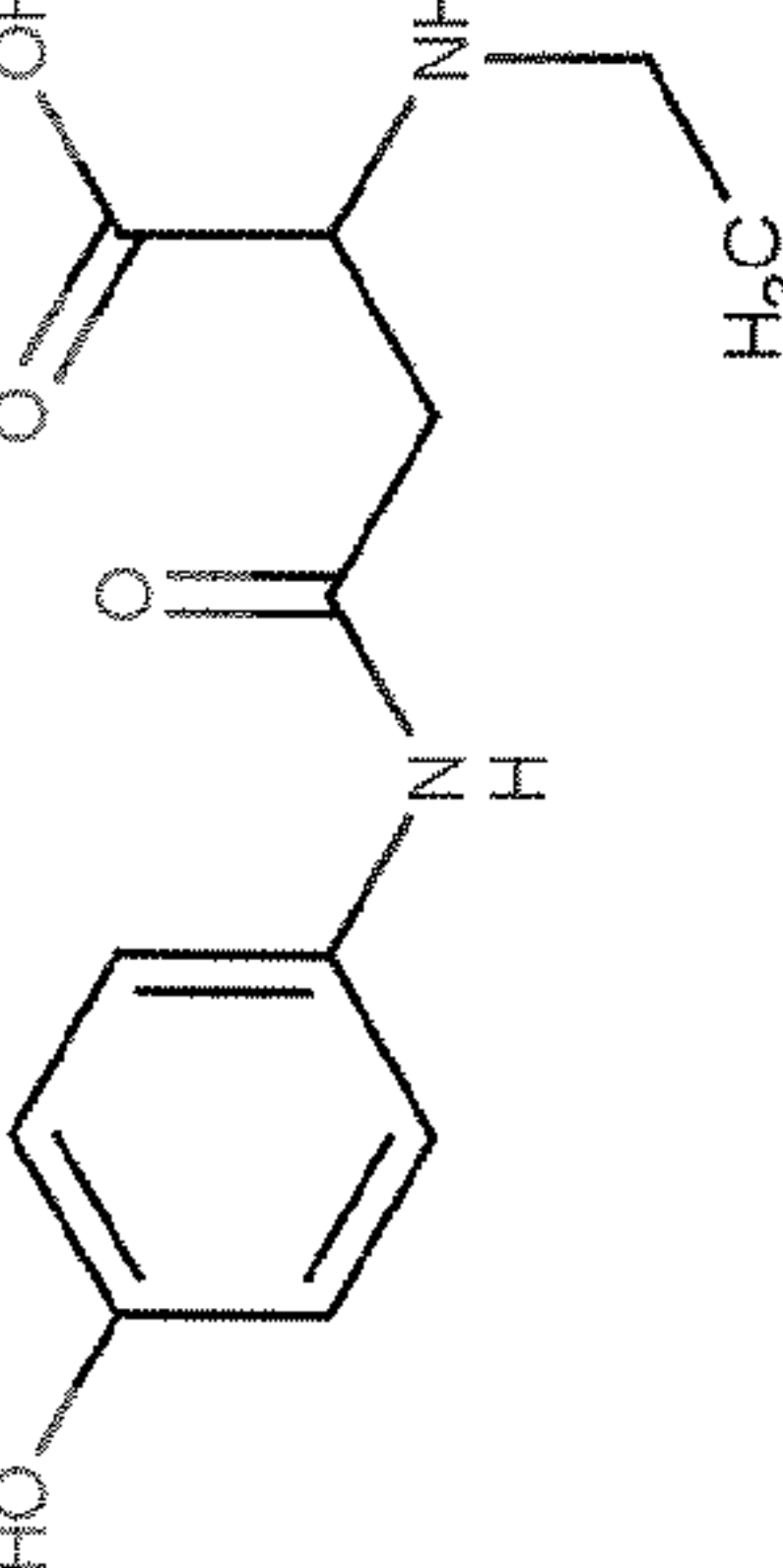
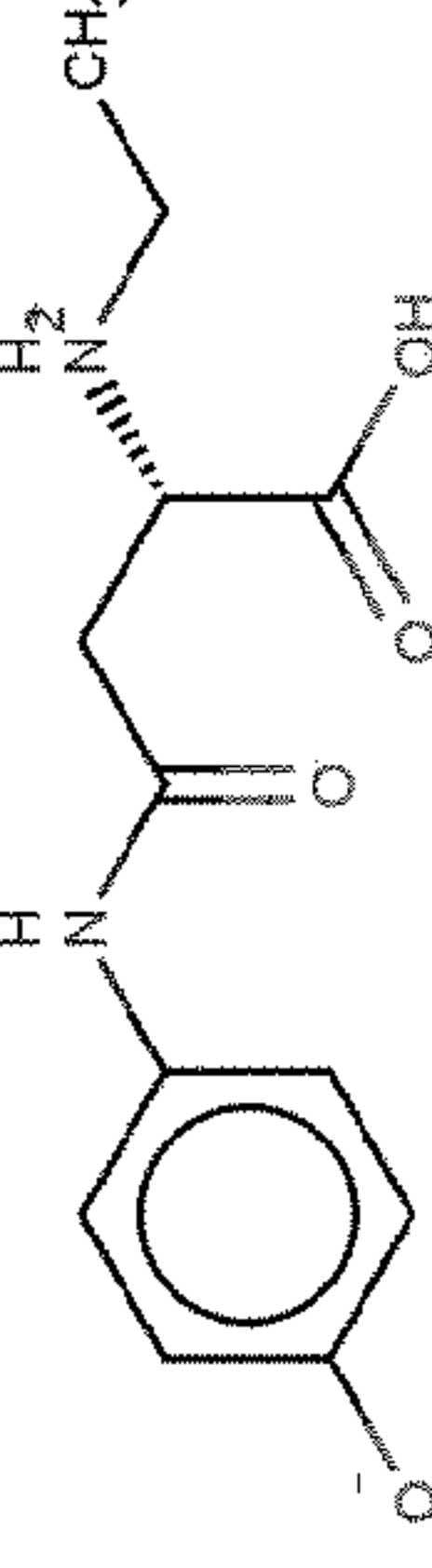
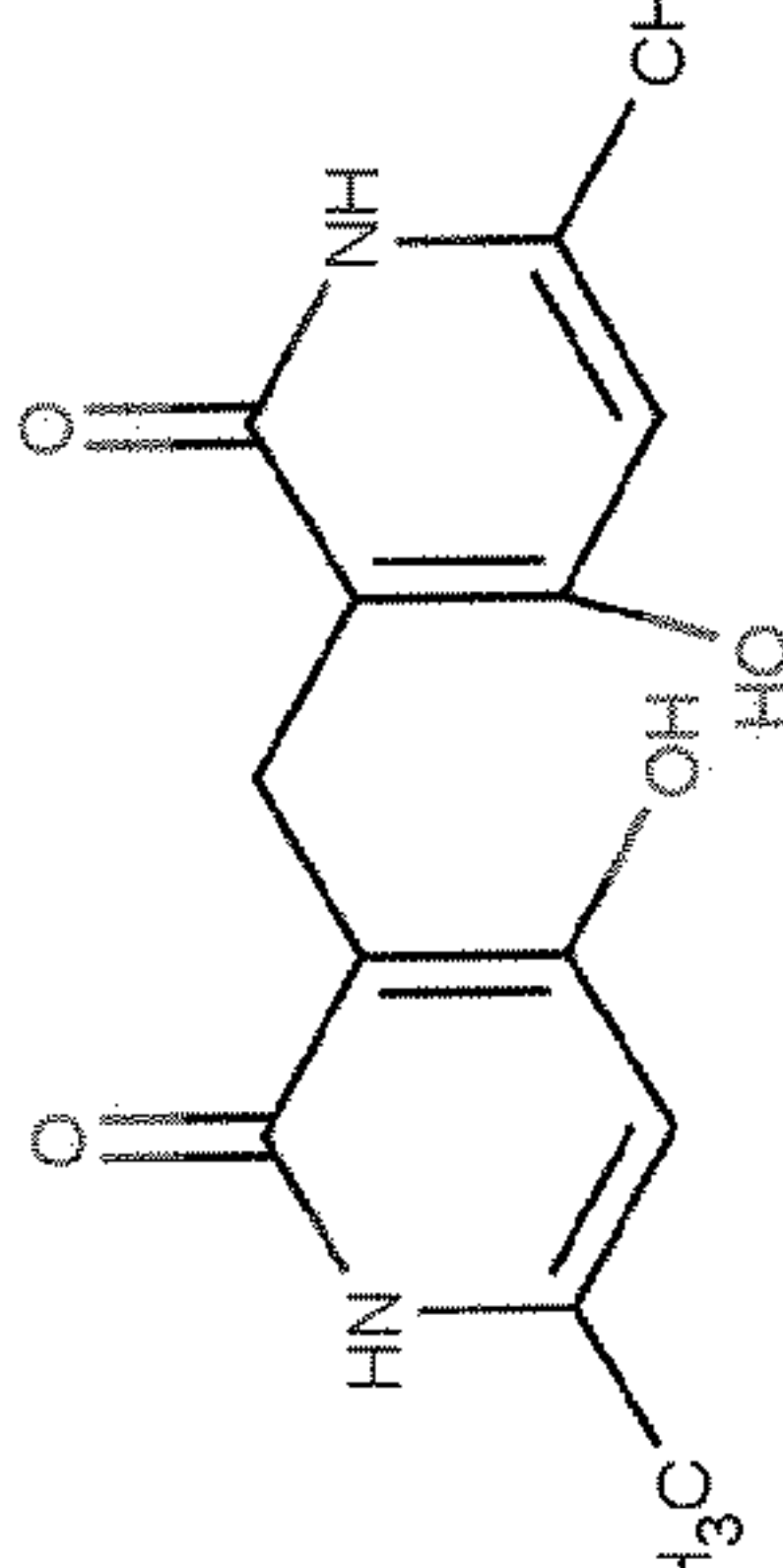
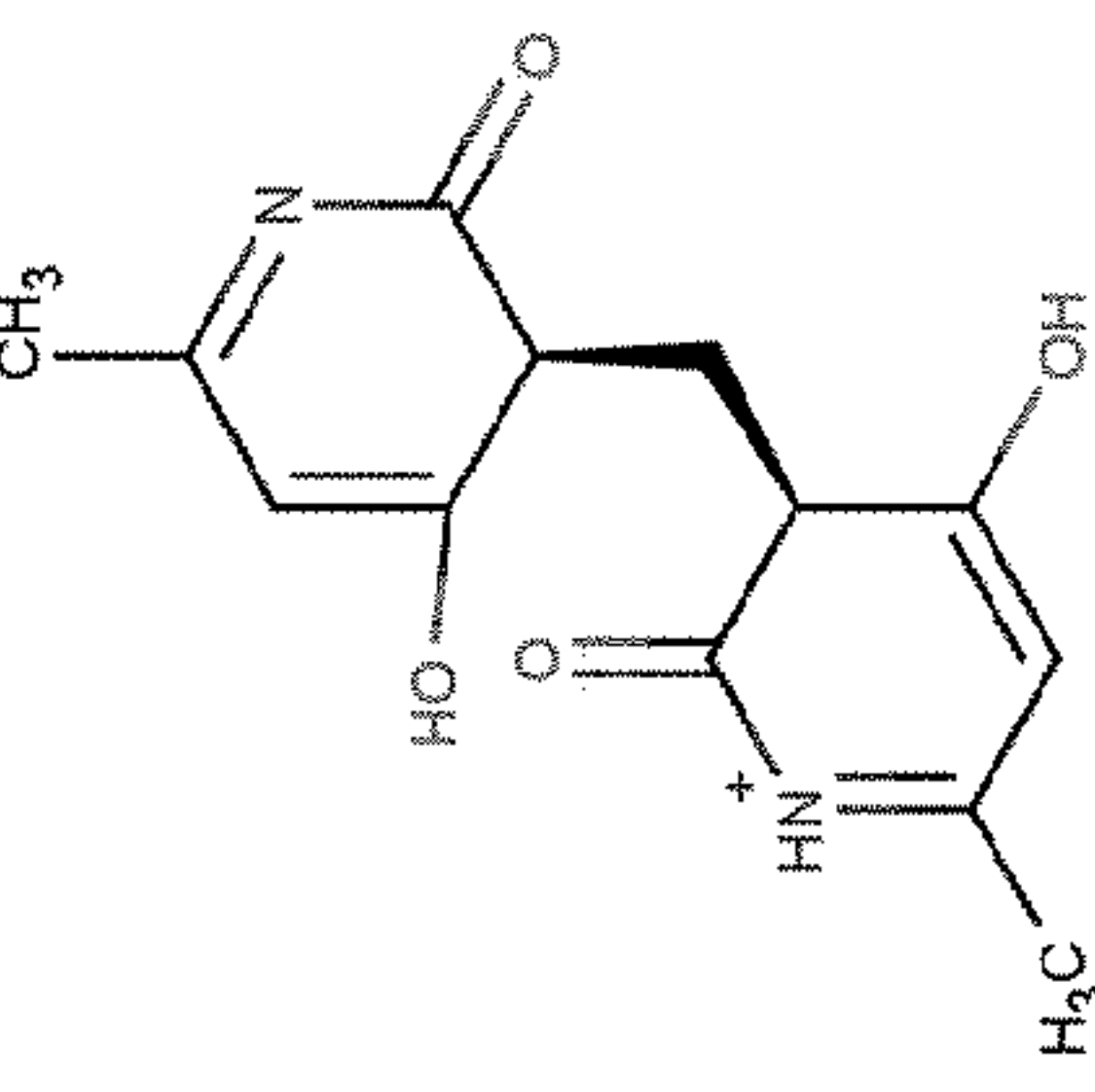
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 127		F2185-0014		F2185-0014_3_2 (1 of 14)
Compound 128		F2189-0334		F2189-0334_5_2 (1 of 14)
Compound 129		F2206-0079		F2206-0079_2_1 (1 of 3)

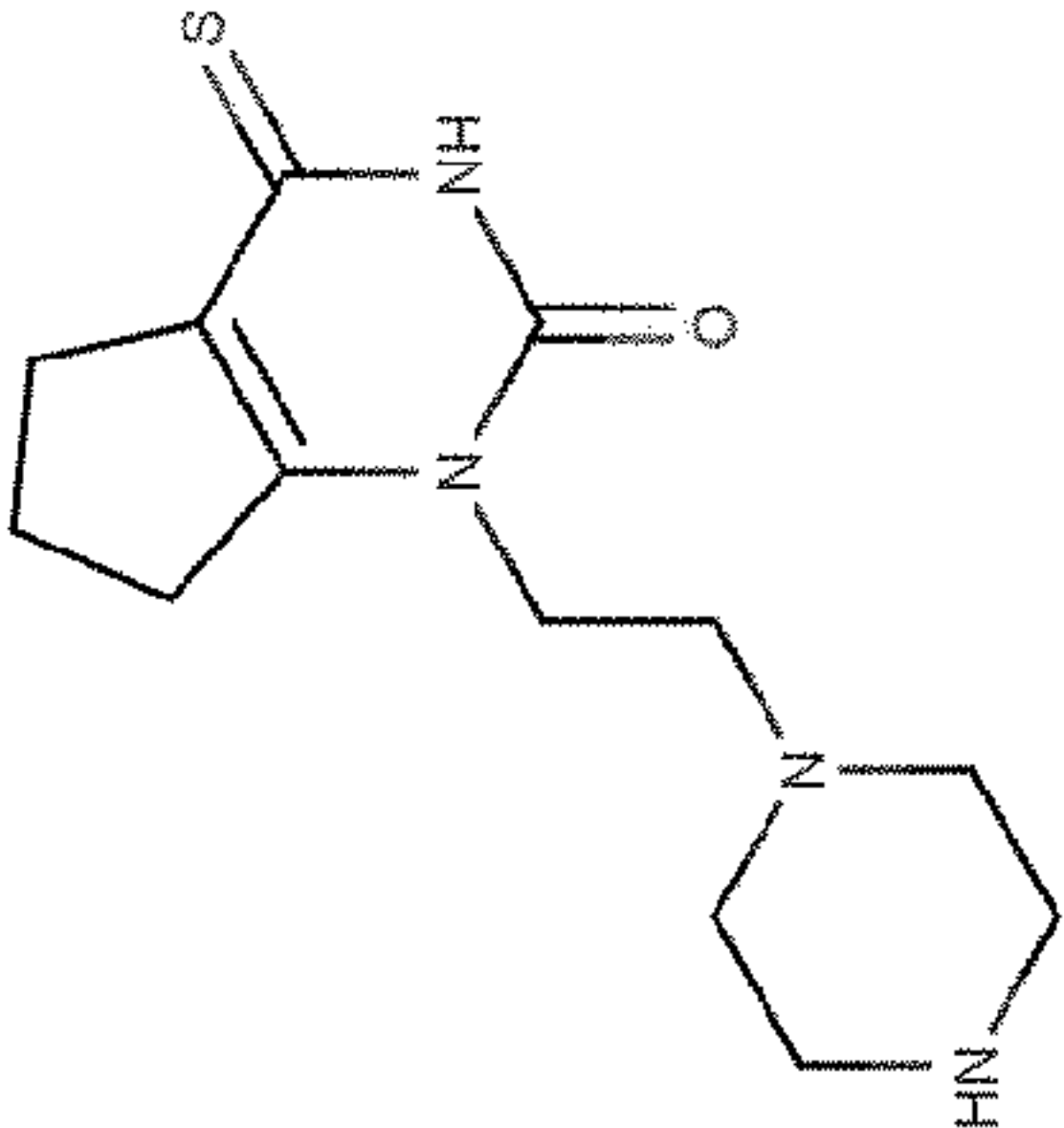
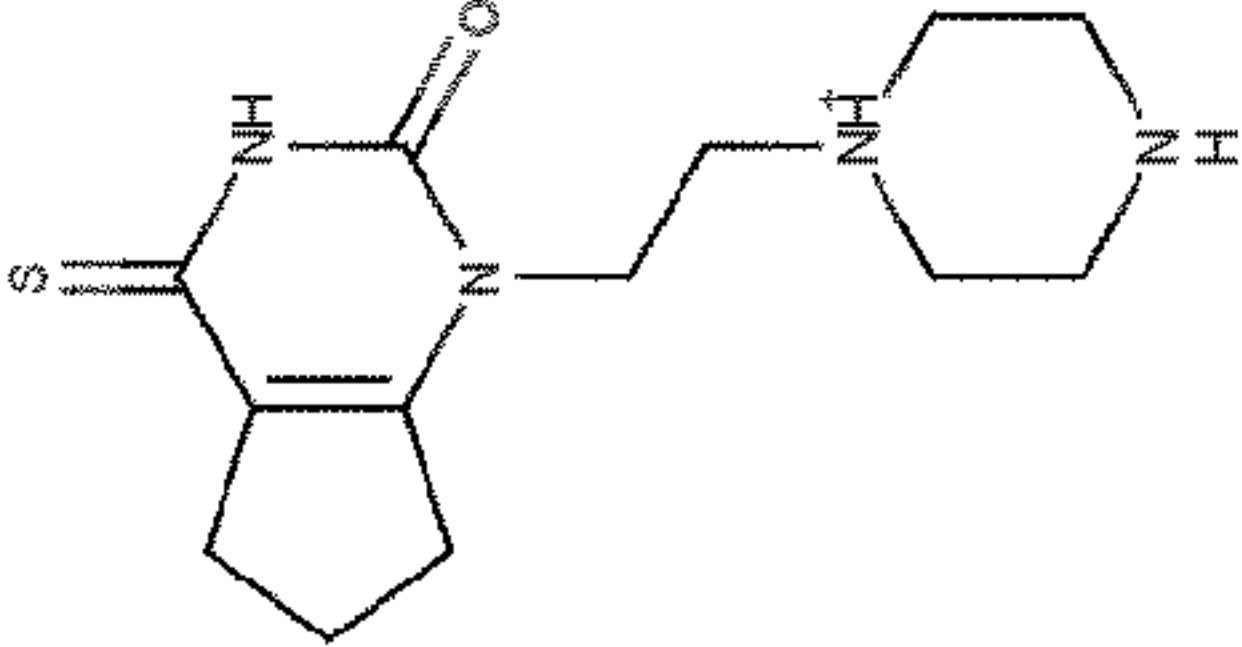
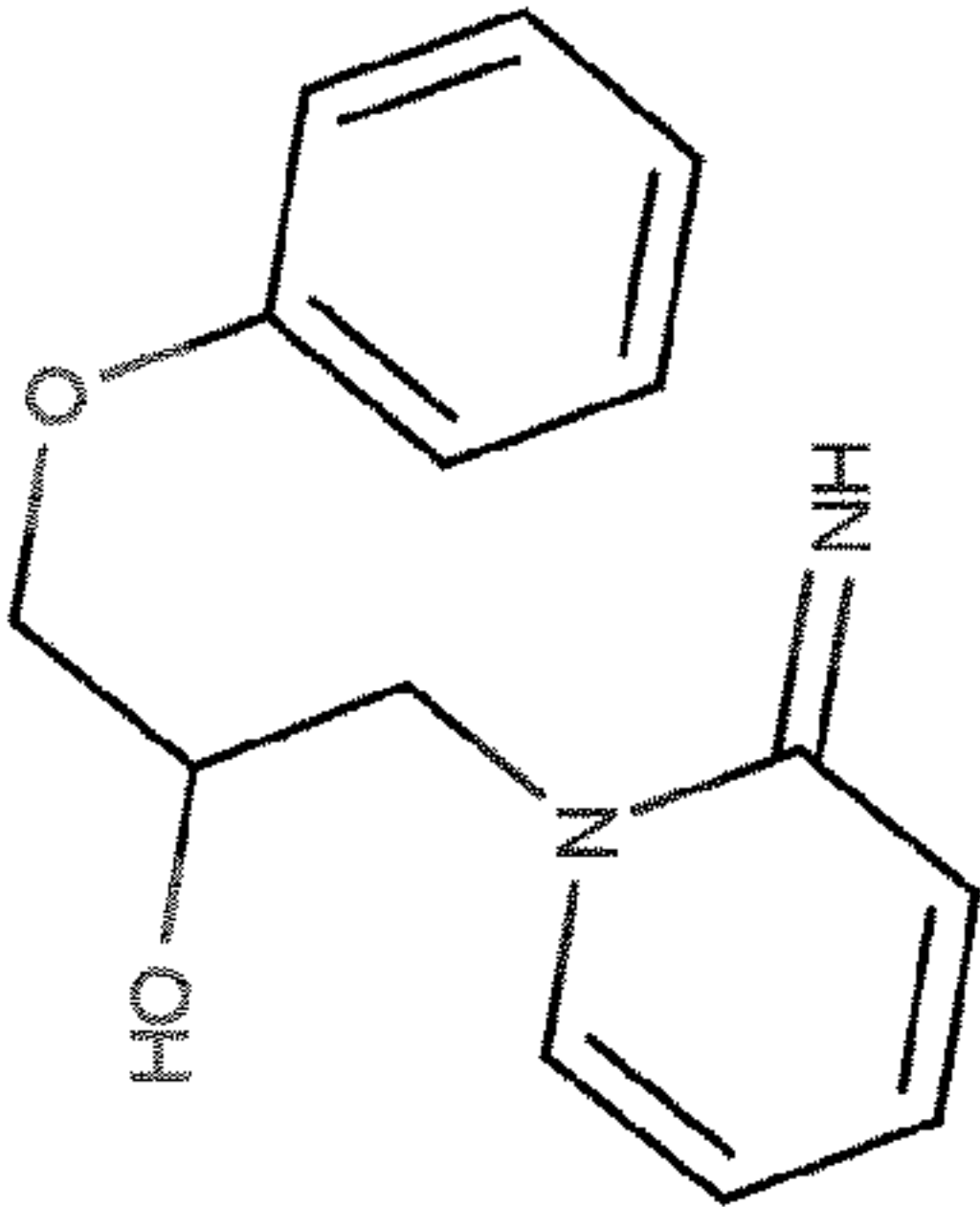
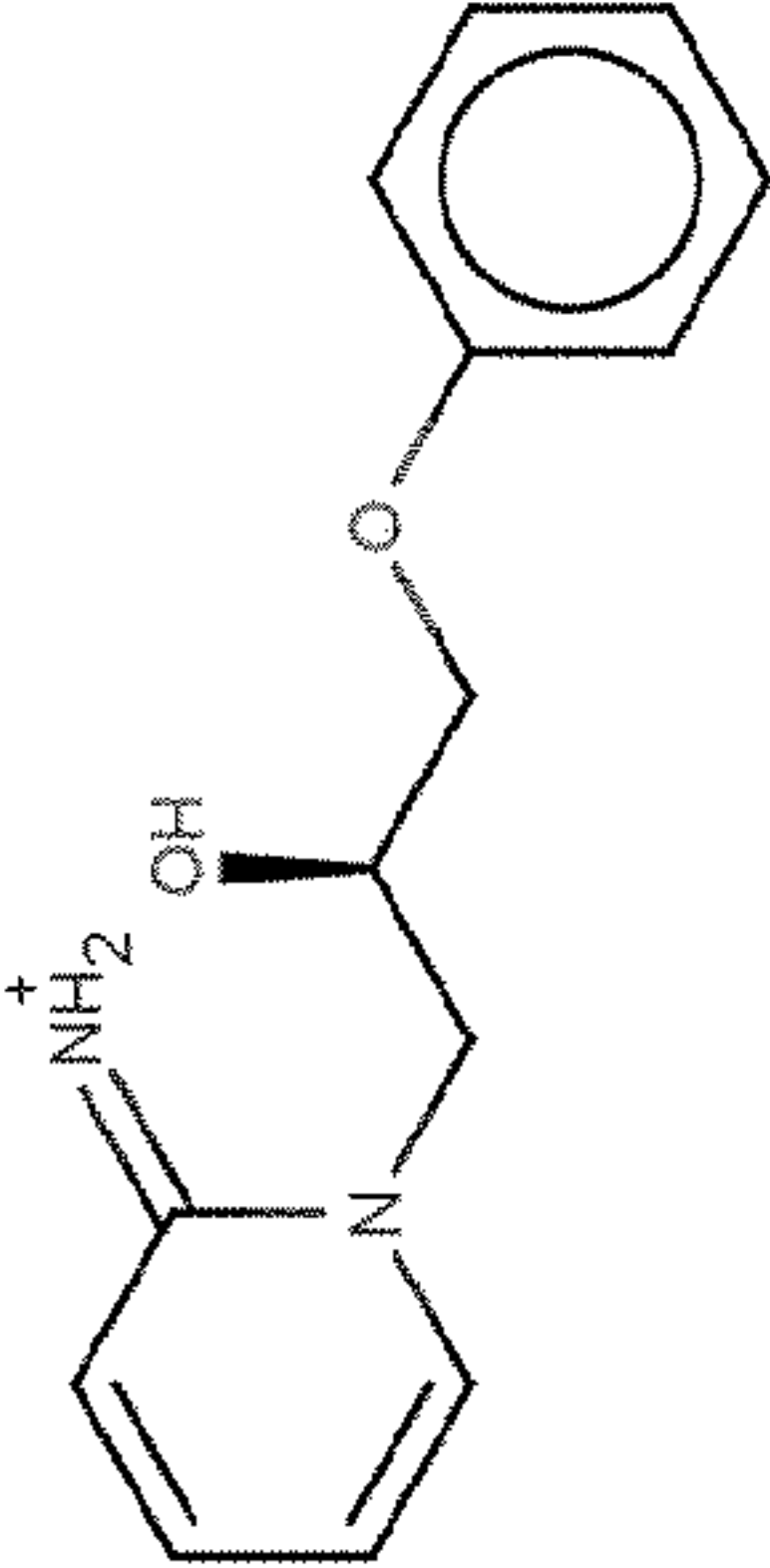
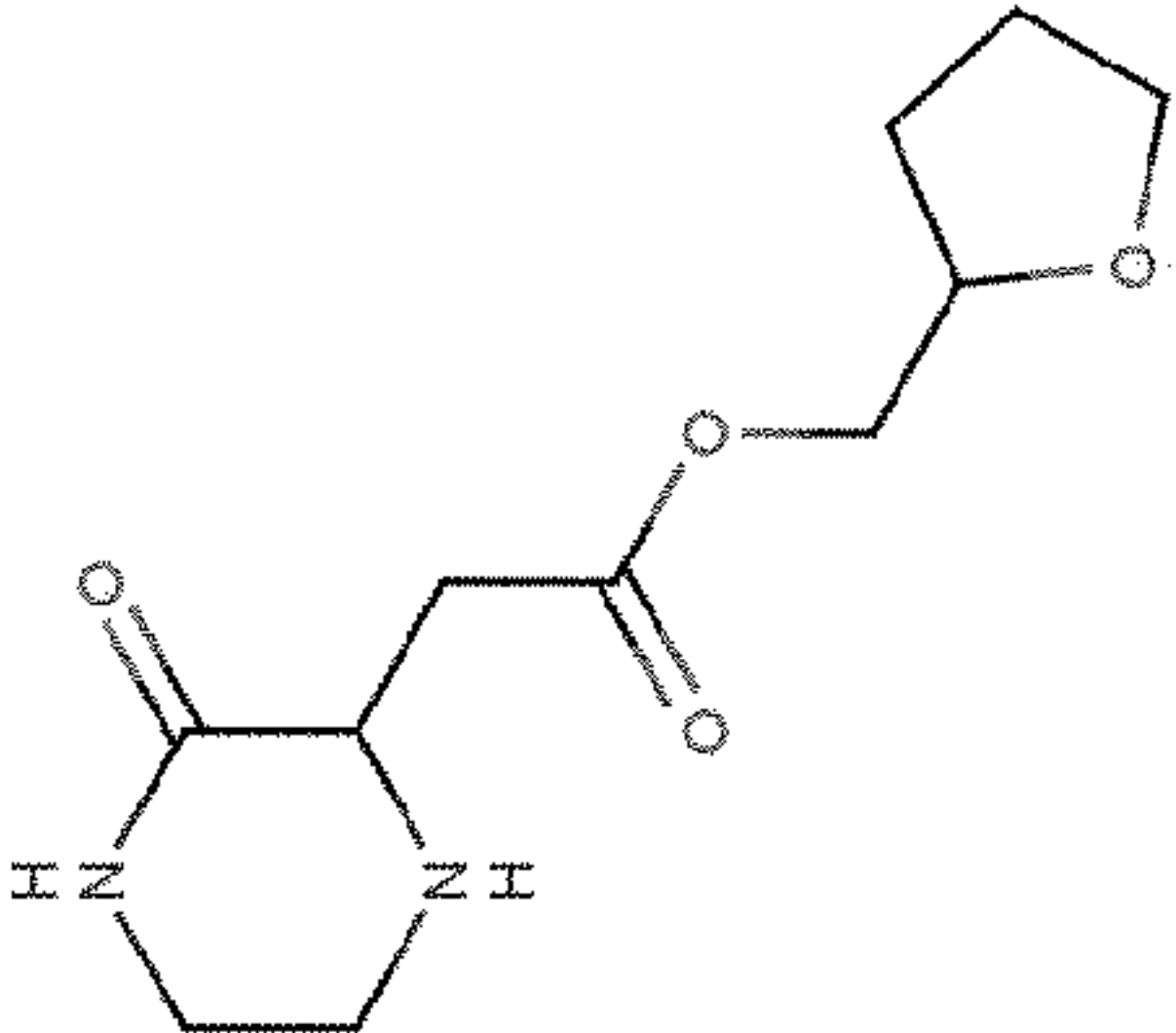
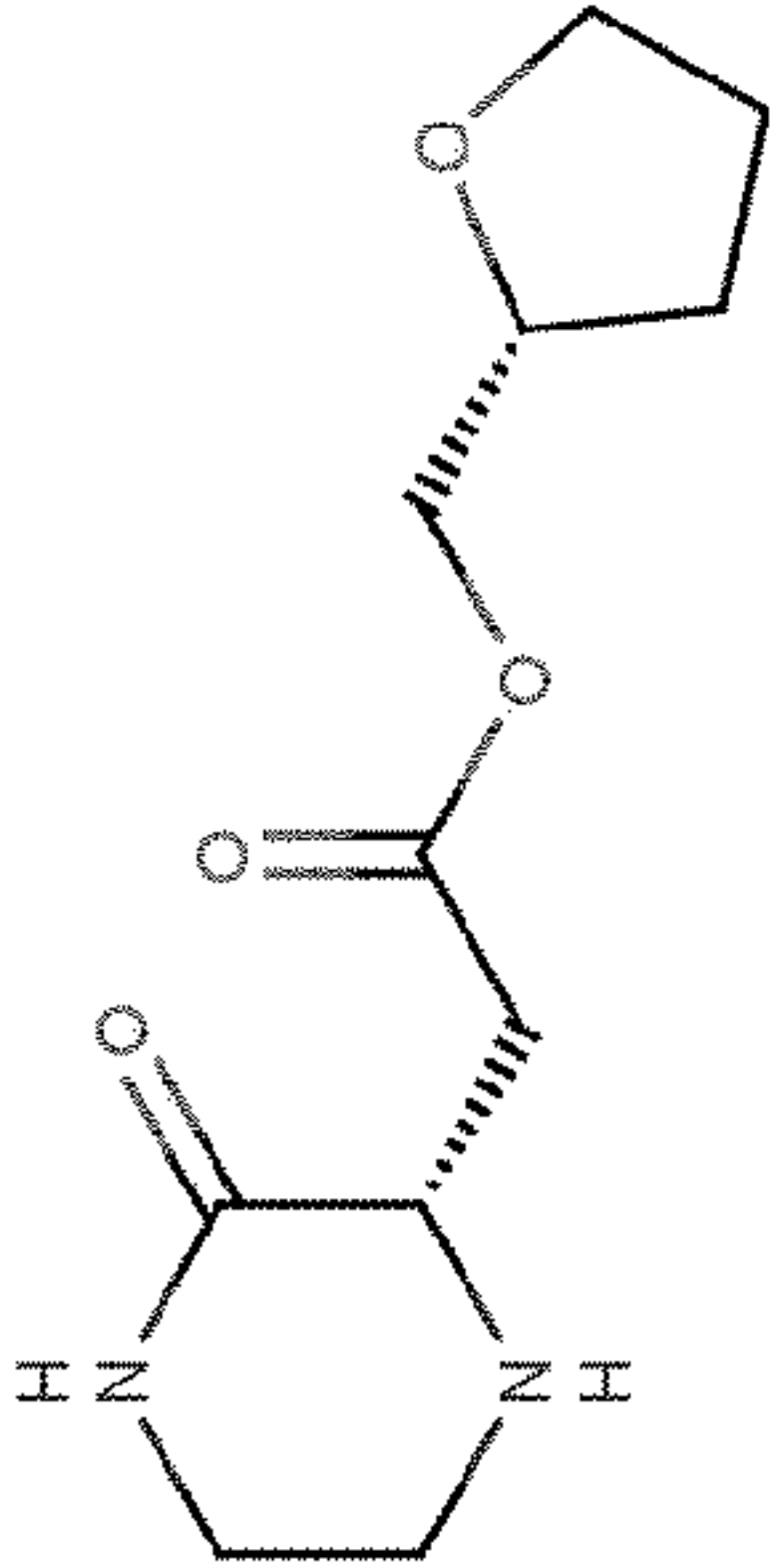
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 130		F2274-0585		F2274-0585_3_1 (1 of 4)
Compound 131		F2416-0083		F2416-0083_1 (1 of 1)
Compound 132		F2503-0105		F2503-0105_22_1 (1 of 39)

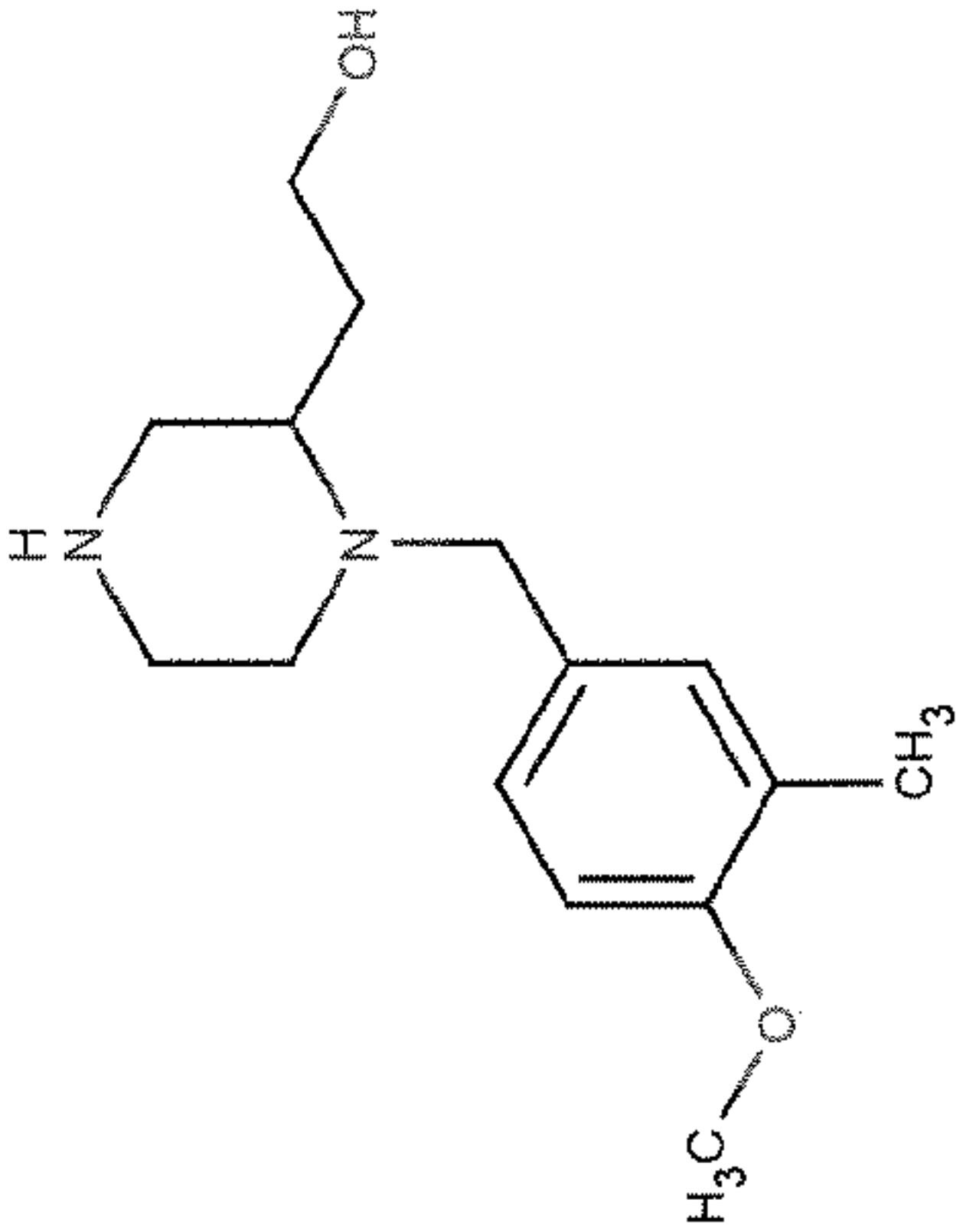
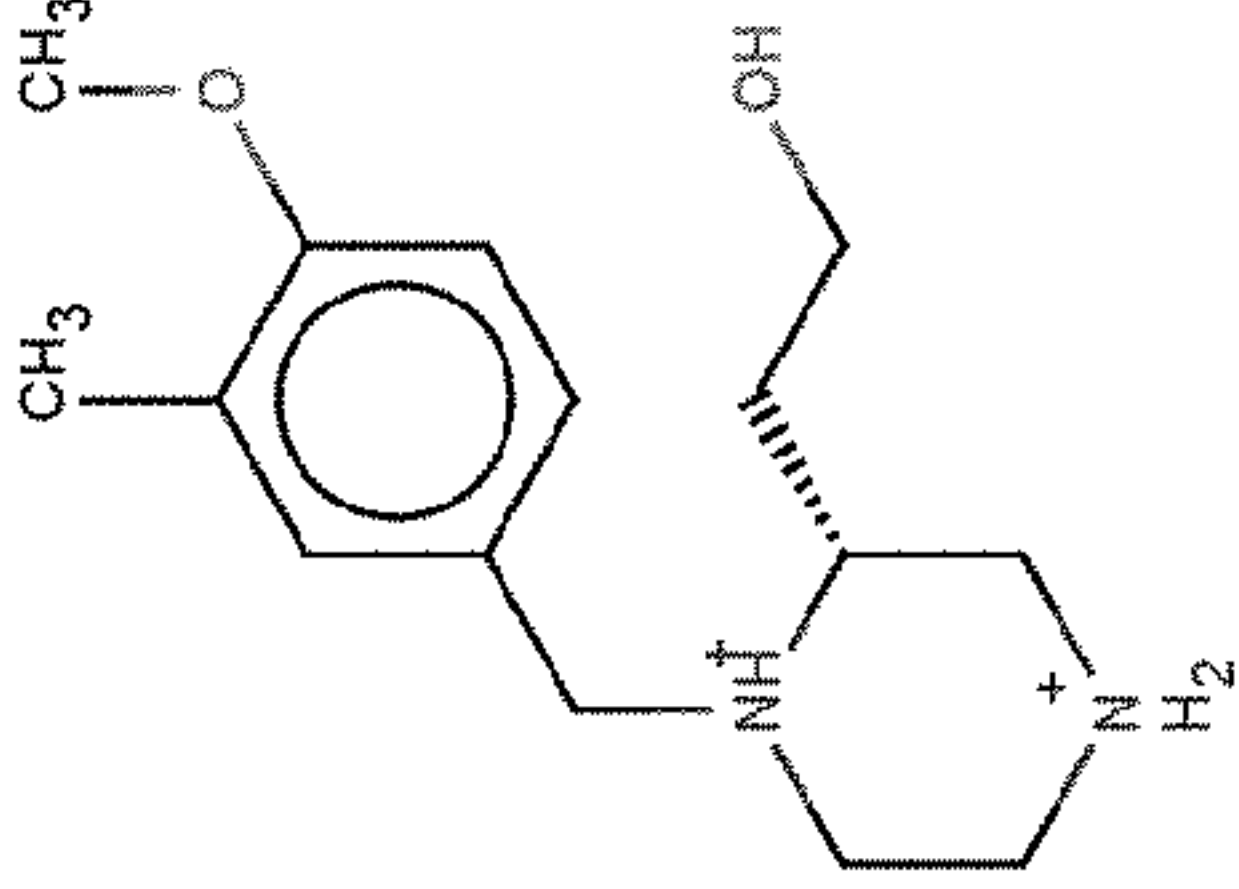
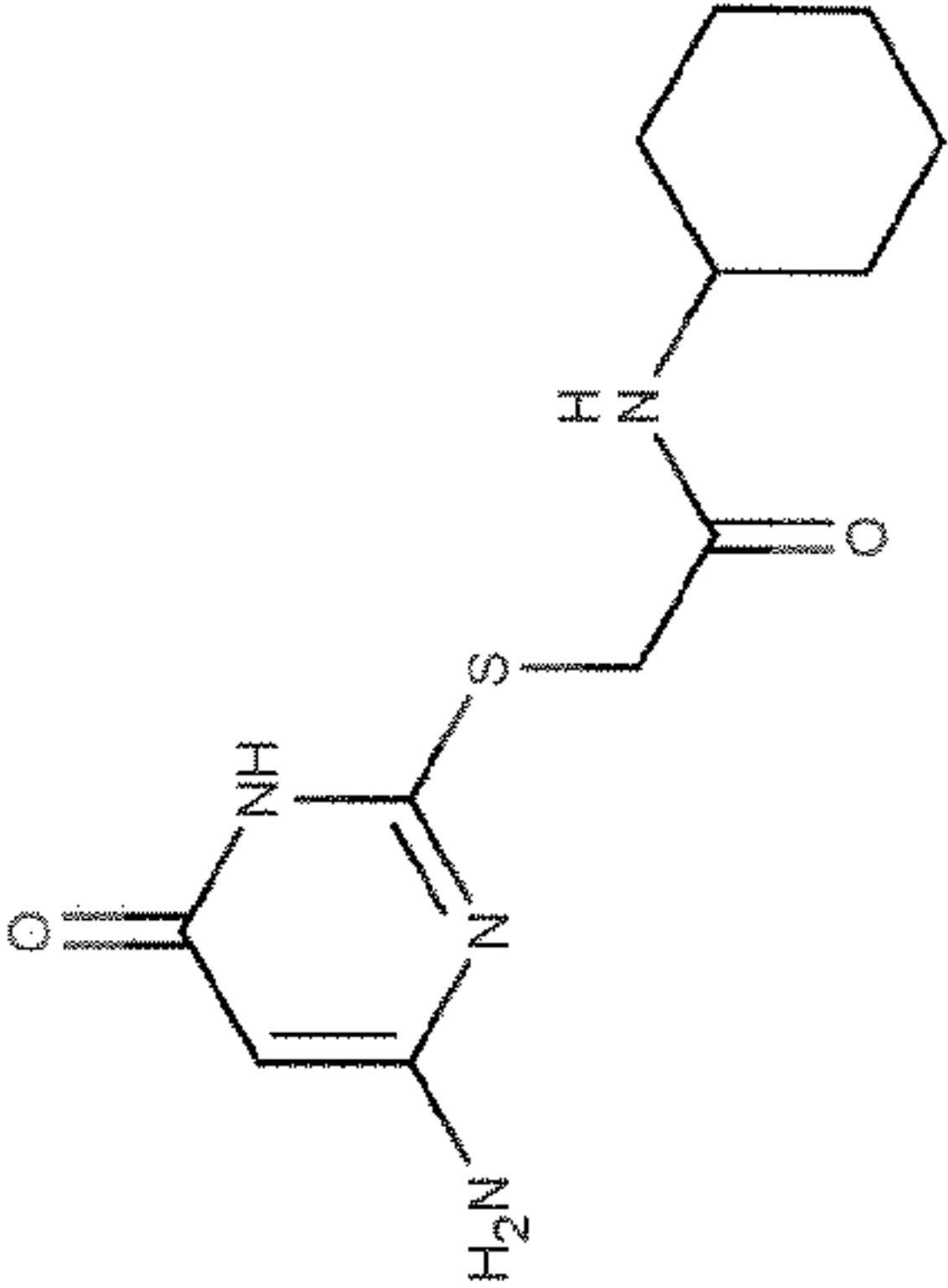
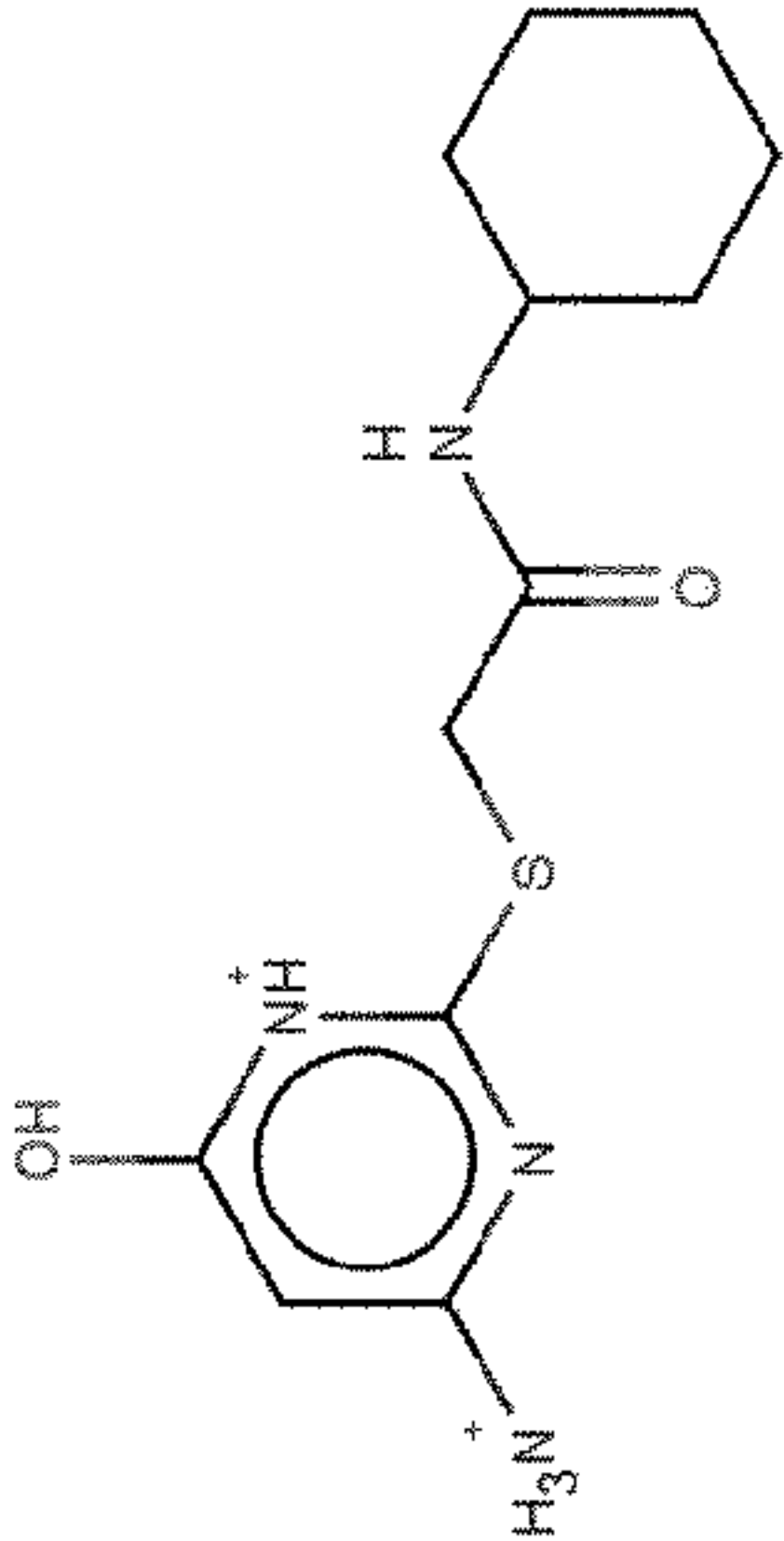
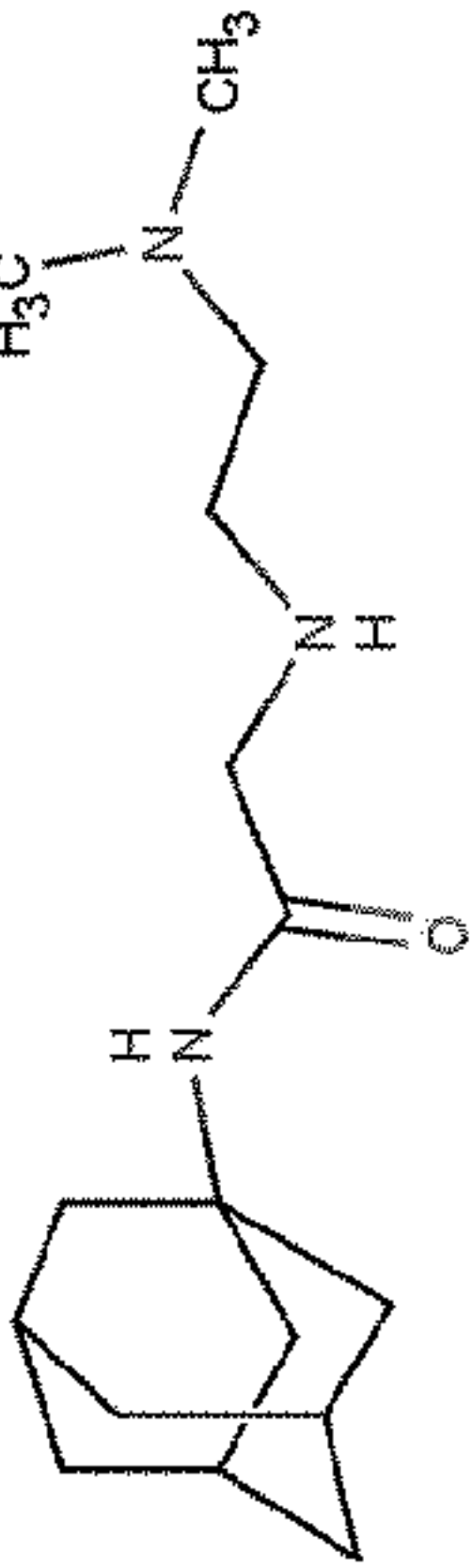
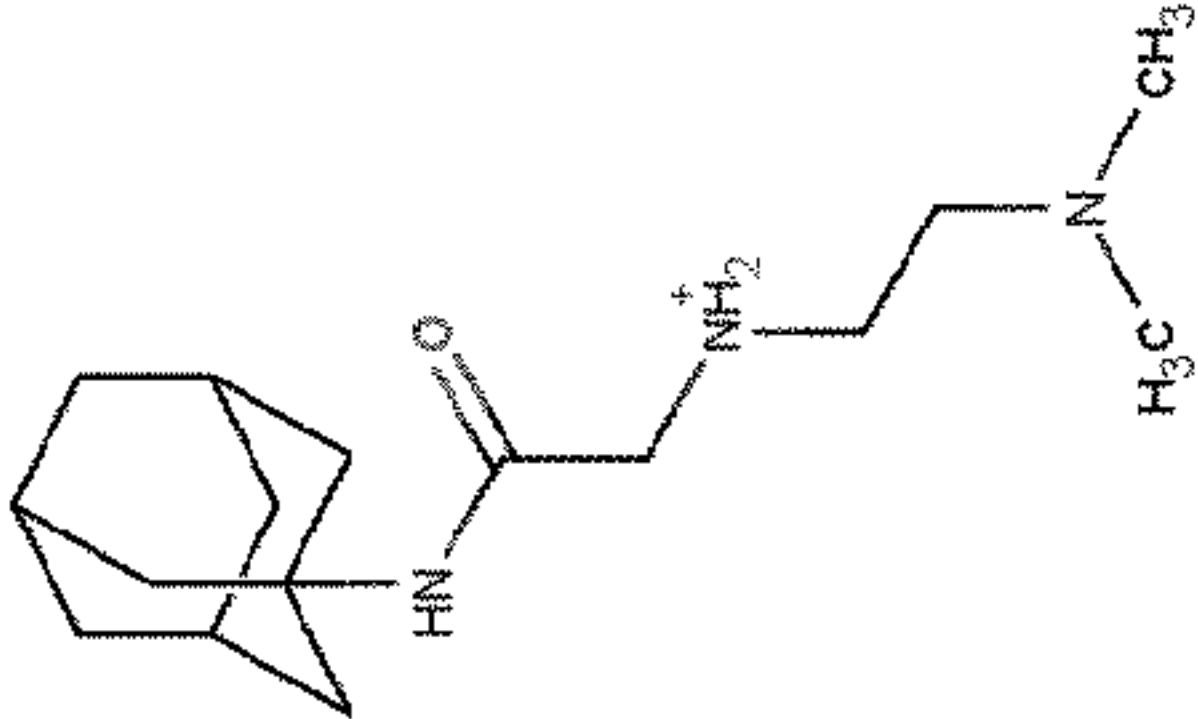
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 133		F2551-0031		F2551-0031_2_1 (1 of 18)
Compound 134		F2622-0072		F2622-0072_41_1 (1 of 129)
Compound 135		F2622-0218		F2622-0218_20_1 (1 of 40)

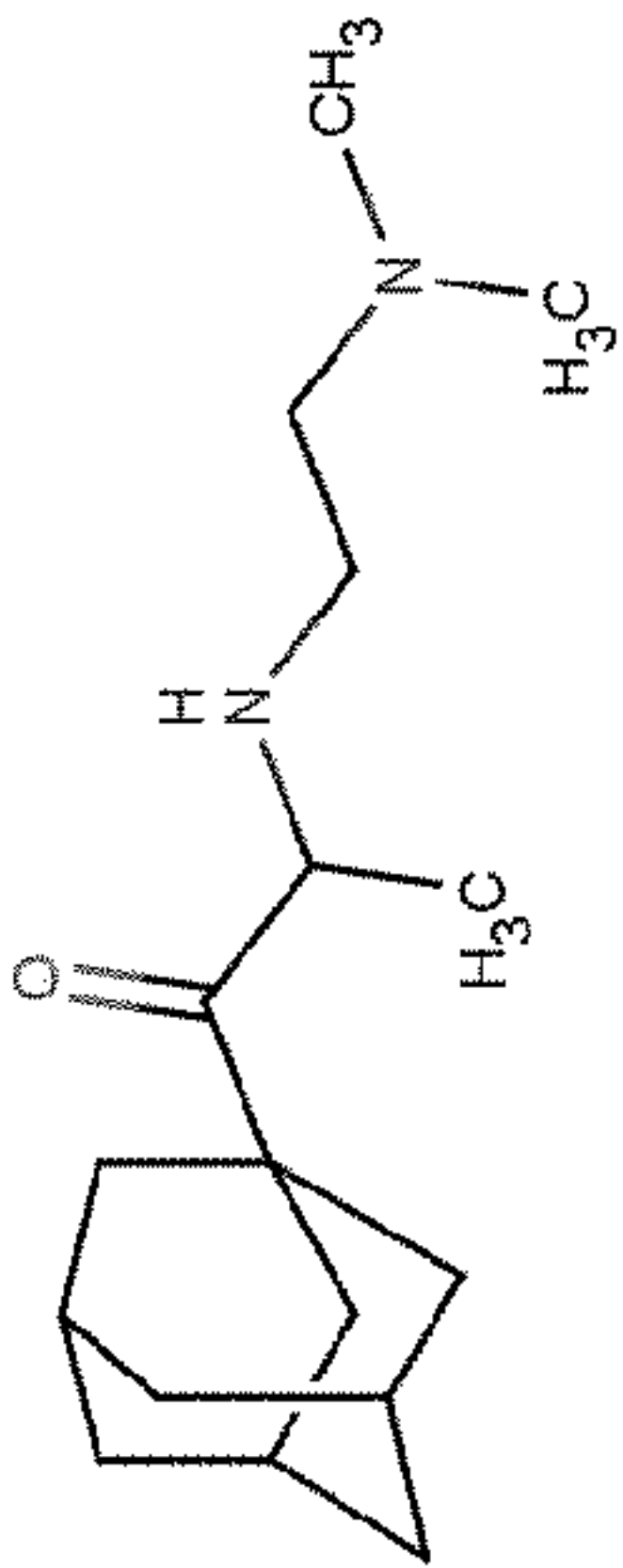
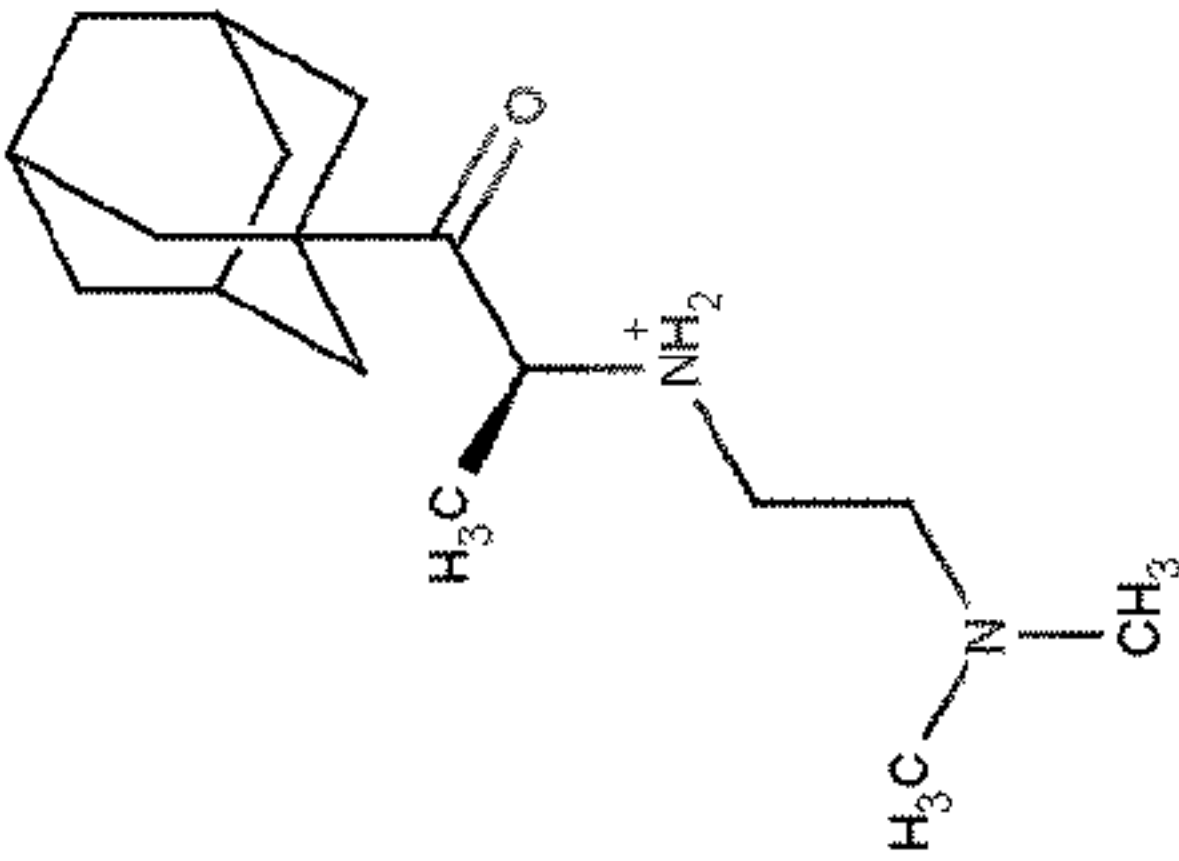
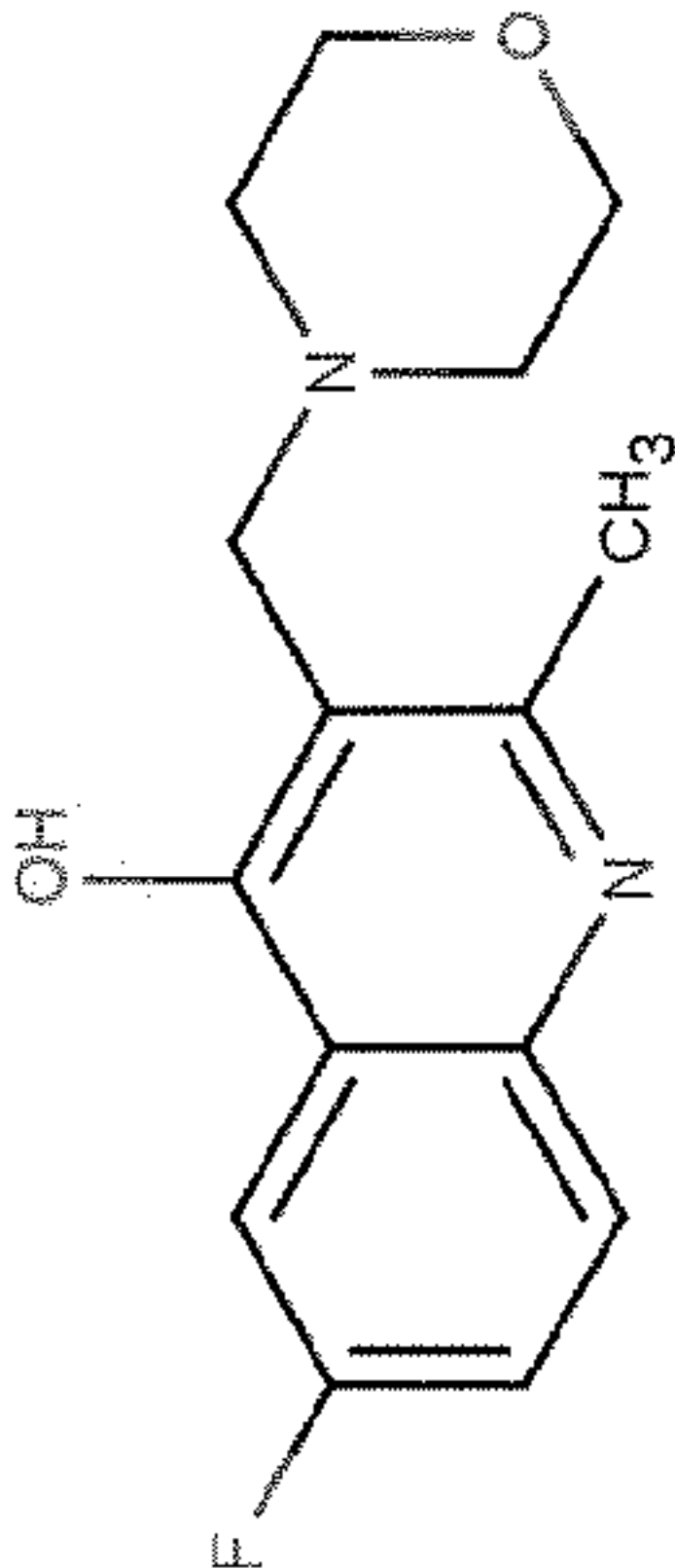
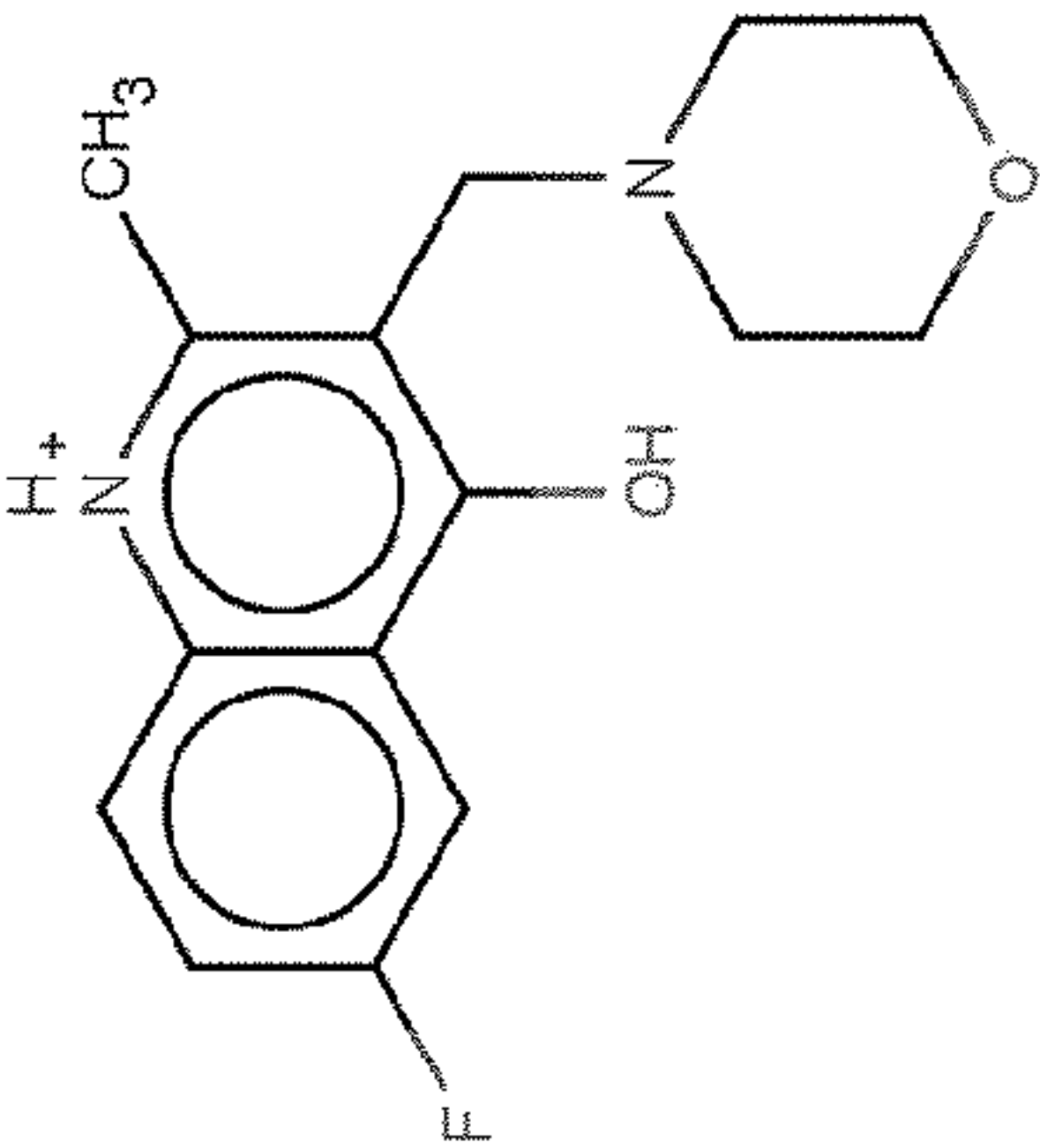
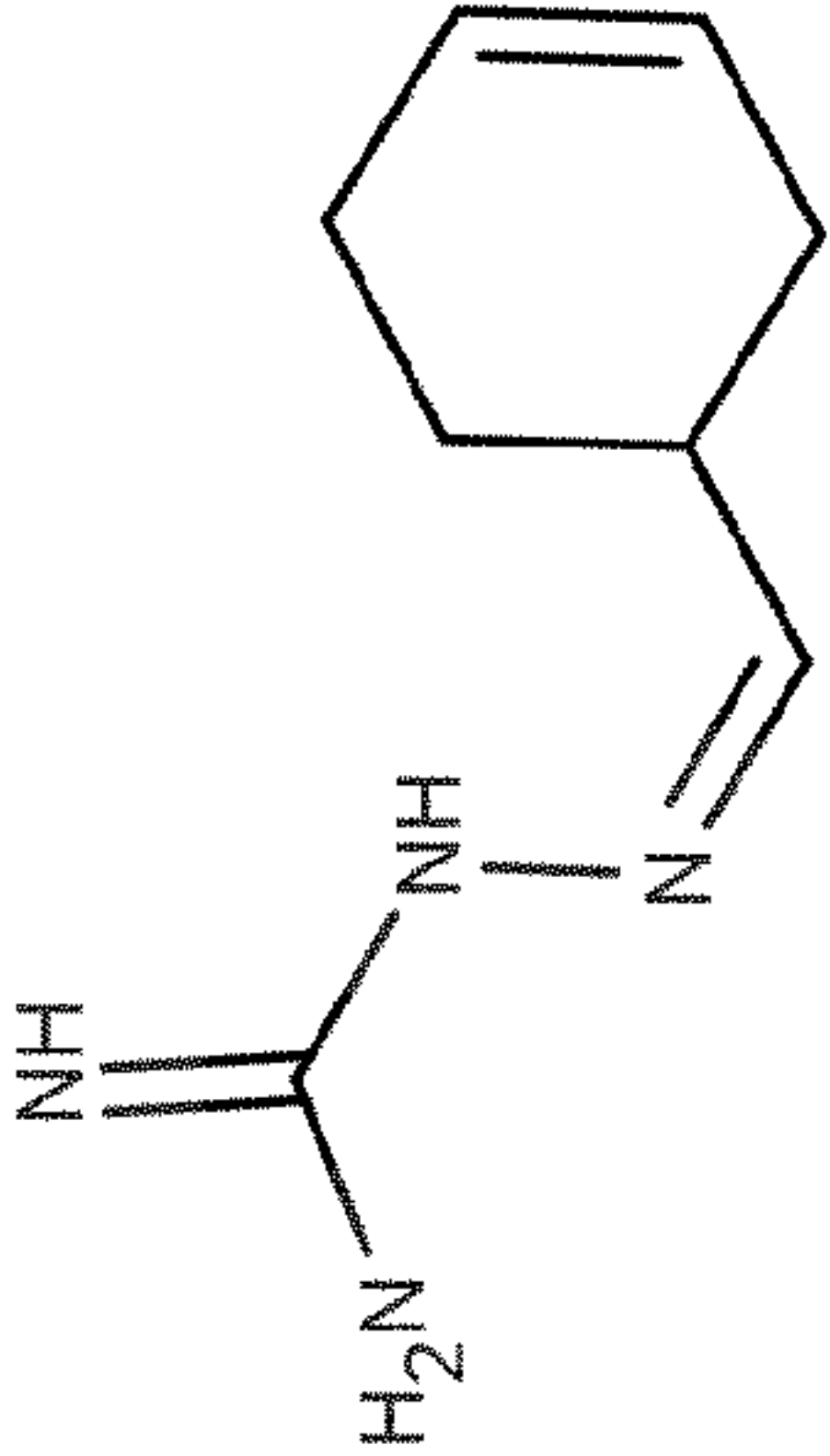
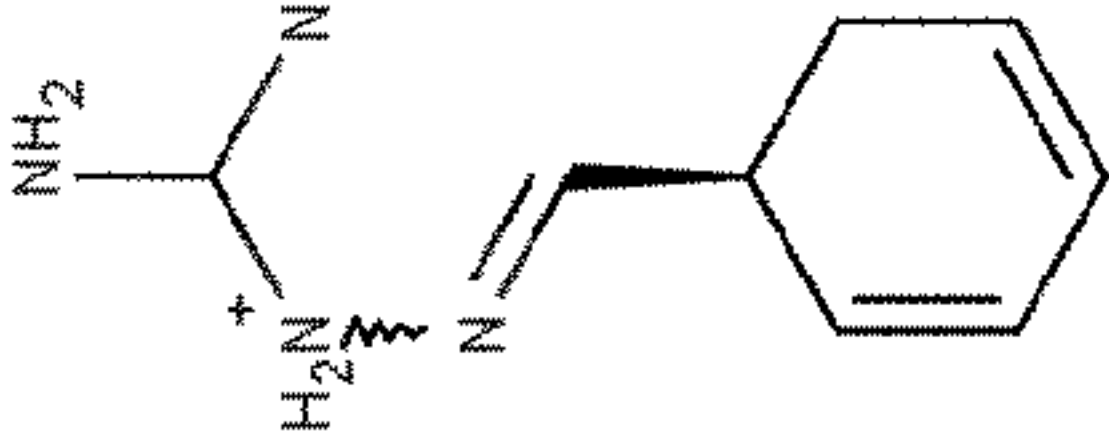
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 136		F2624-0002		F2624-0002_7_1 (1 of 40)
Compound 137		F2624-0008		F2624-0008_7_1 (1 of 40)
Compound 138		F2643-0132		F2643-0132_2_1 (1 of 4)

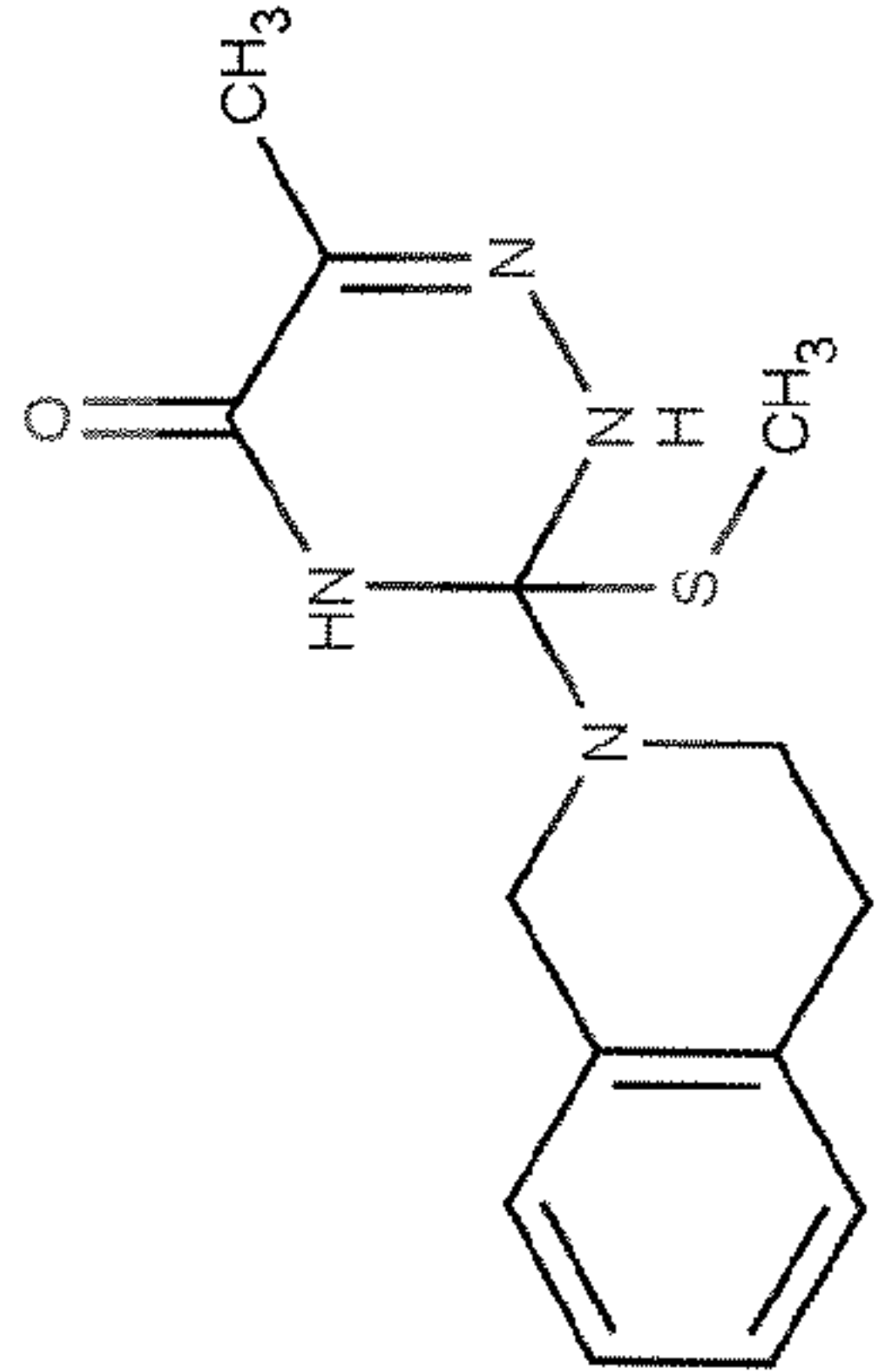
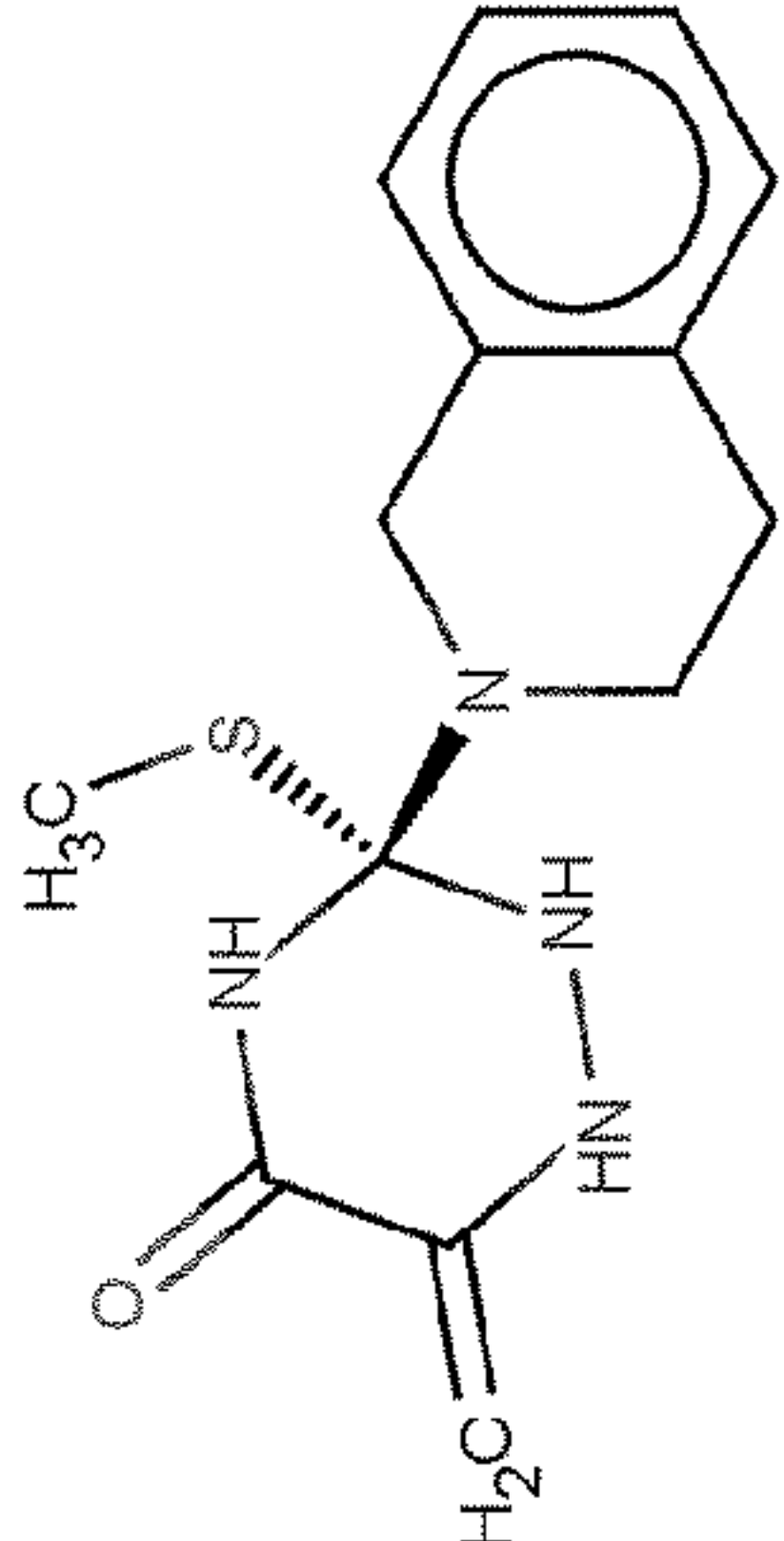
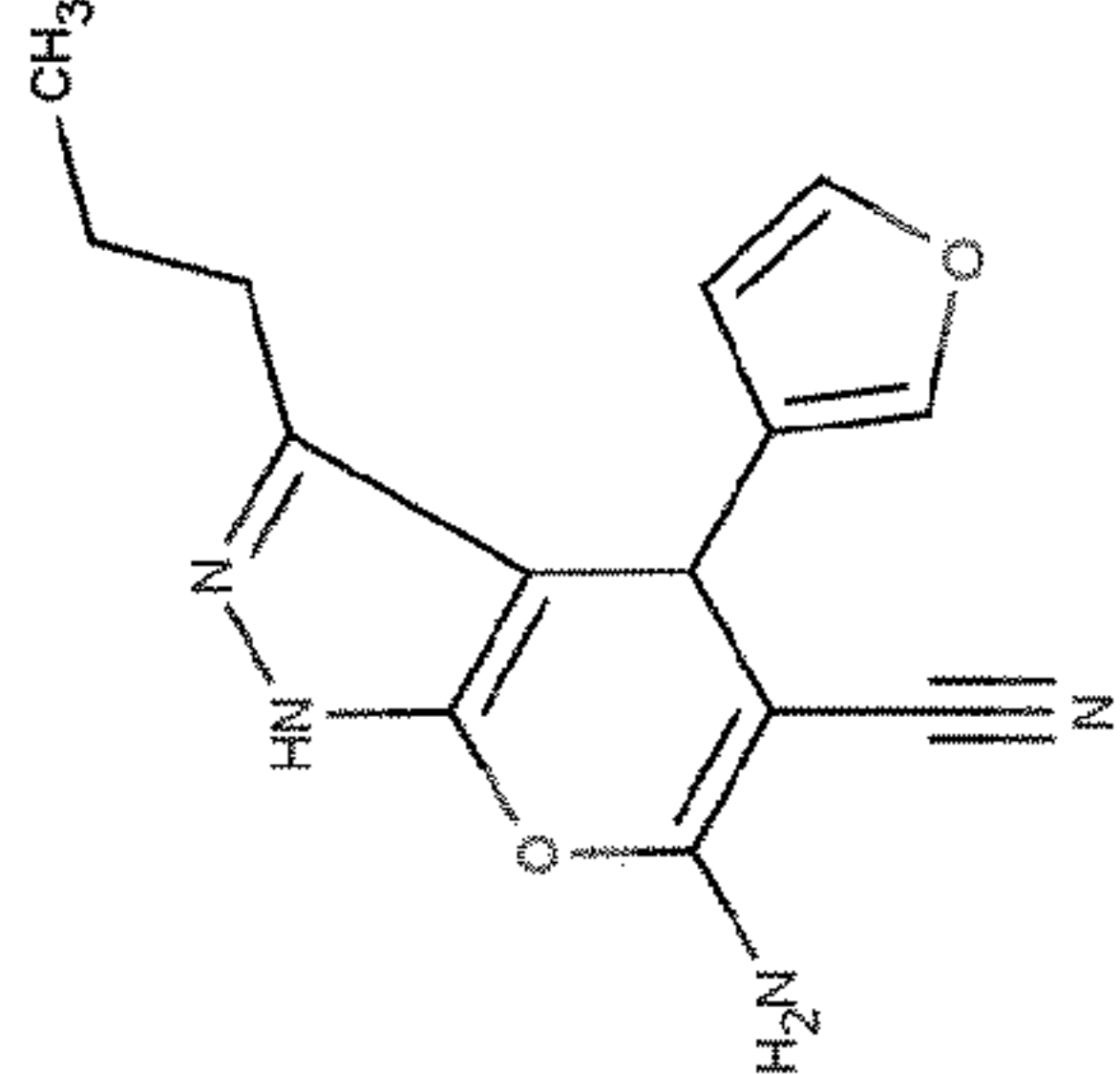
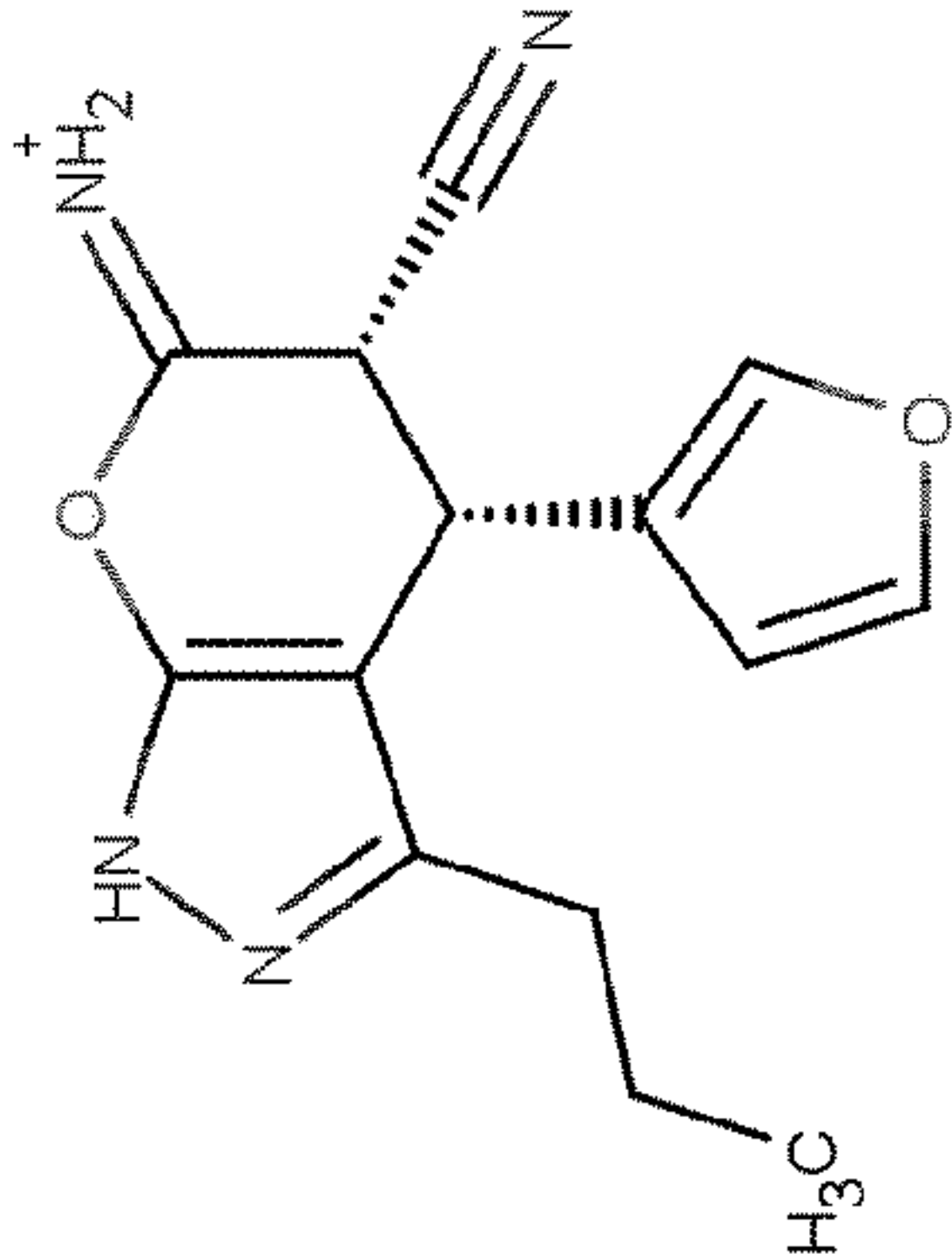
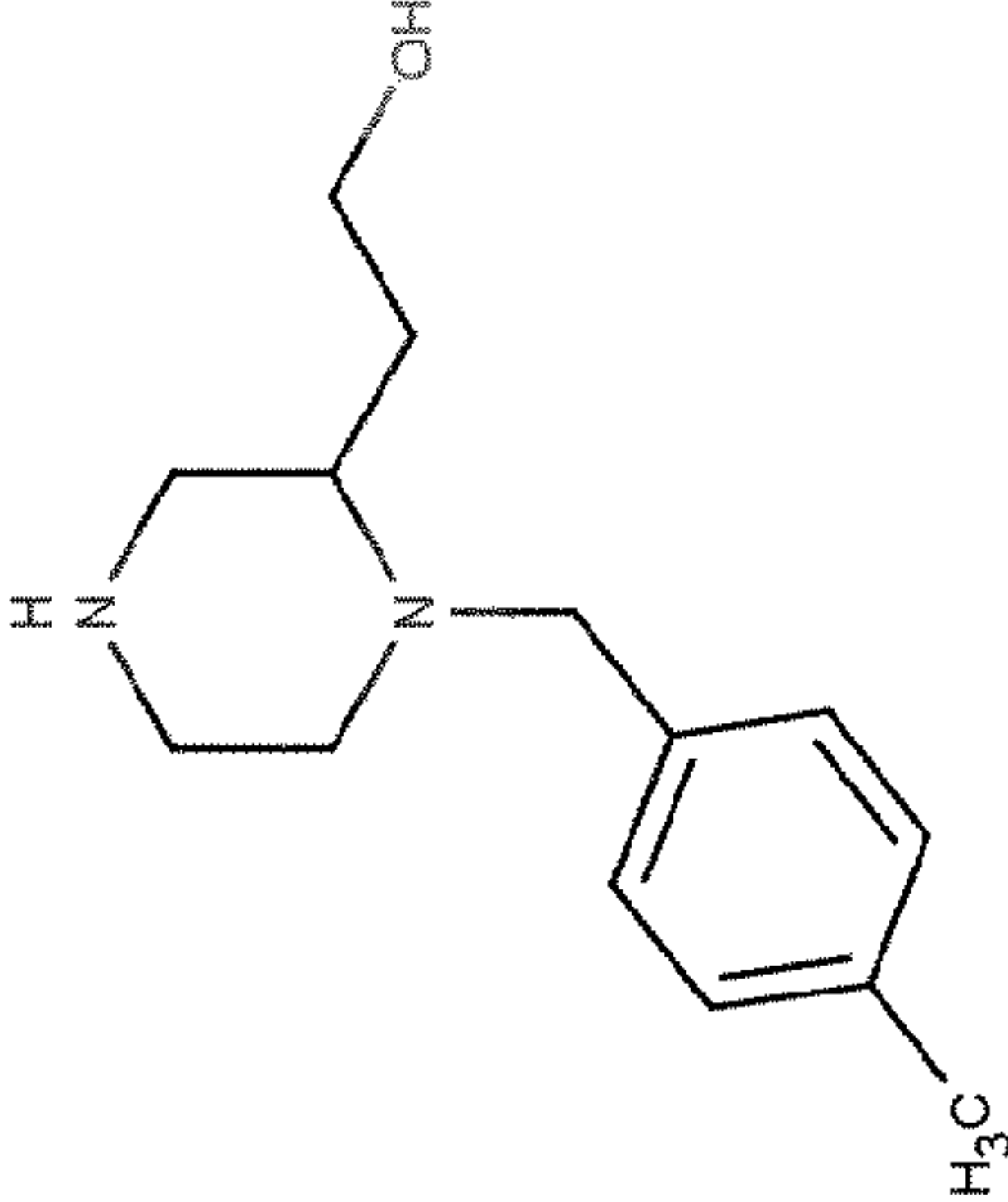
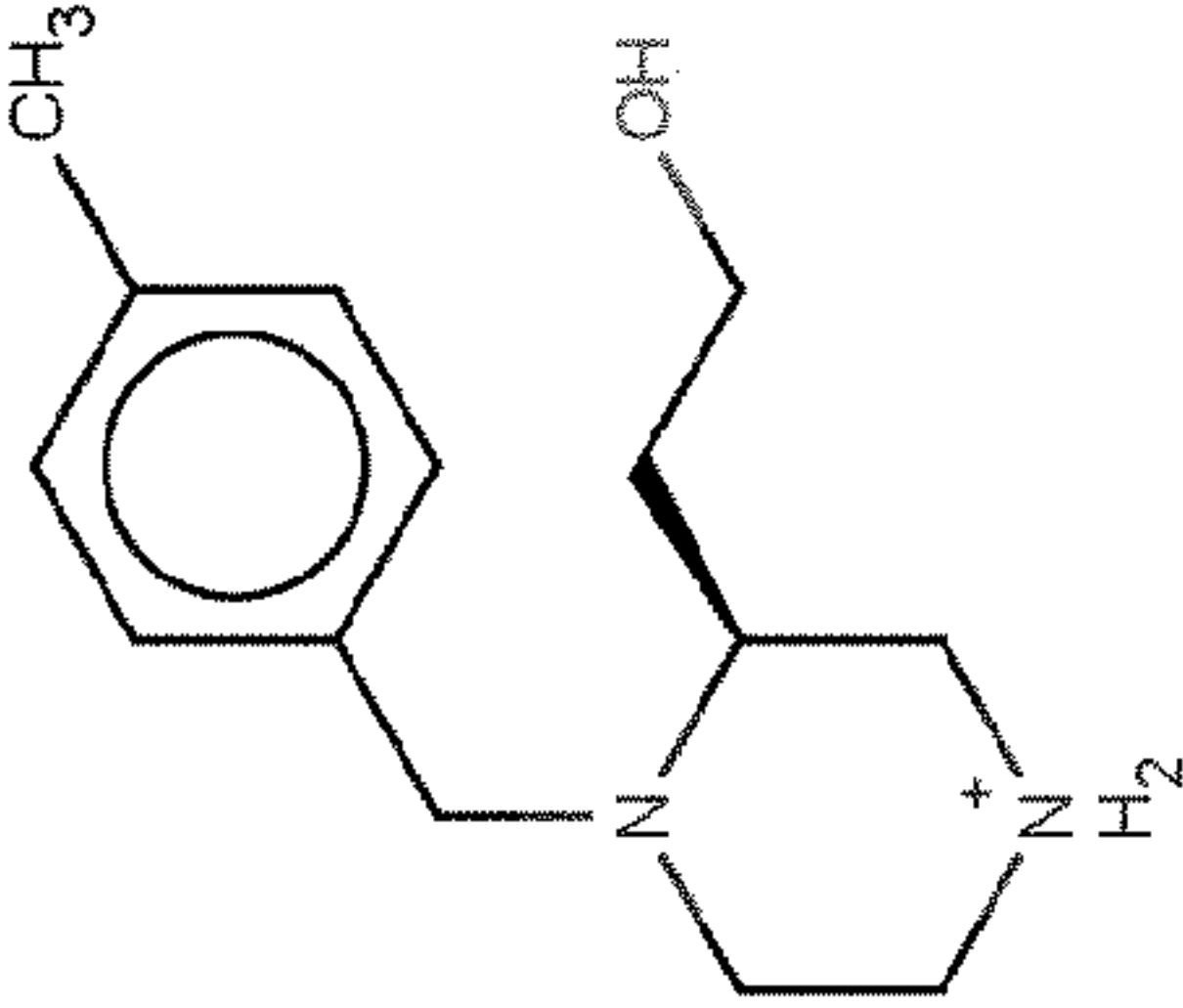
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 139		F2711-1093		F2711-1093_10_1 (1 of 60)
Compound 140		F2713-0139		F2713-0139_4_1 (1 of 5)
Compound 141		F2716-0876		F2716-0876_15_1 (1 of 14)

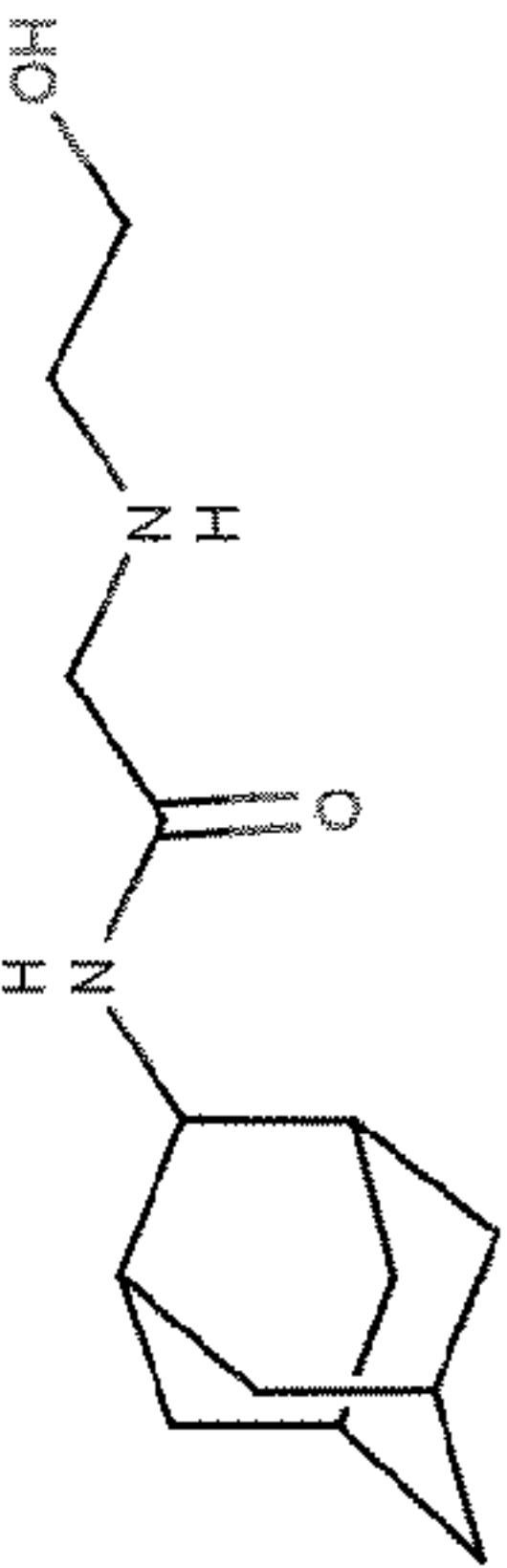
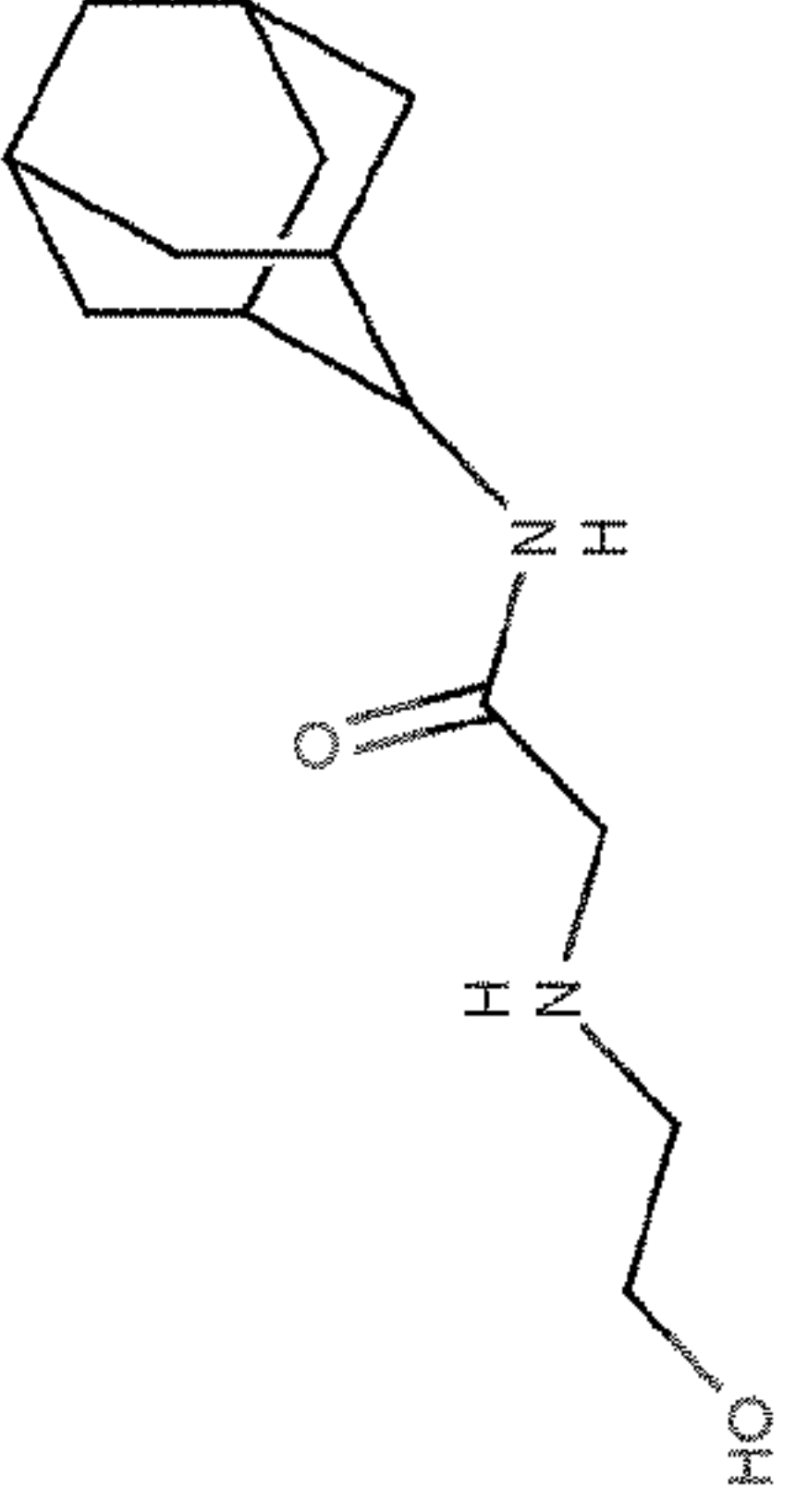
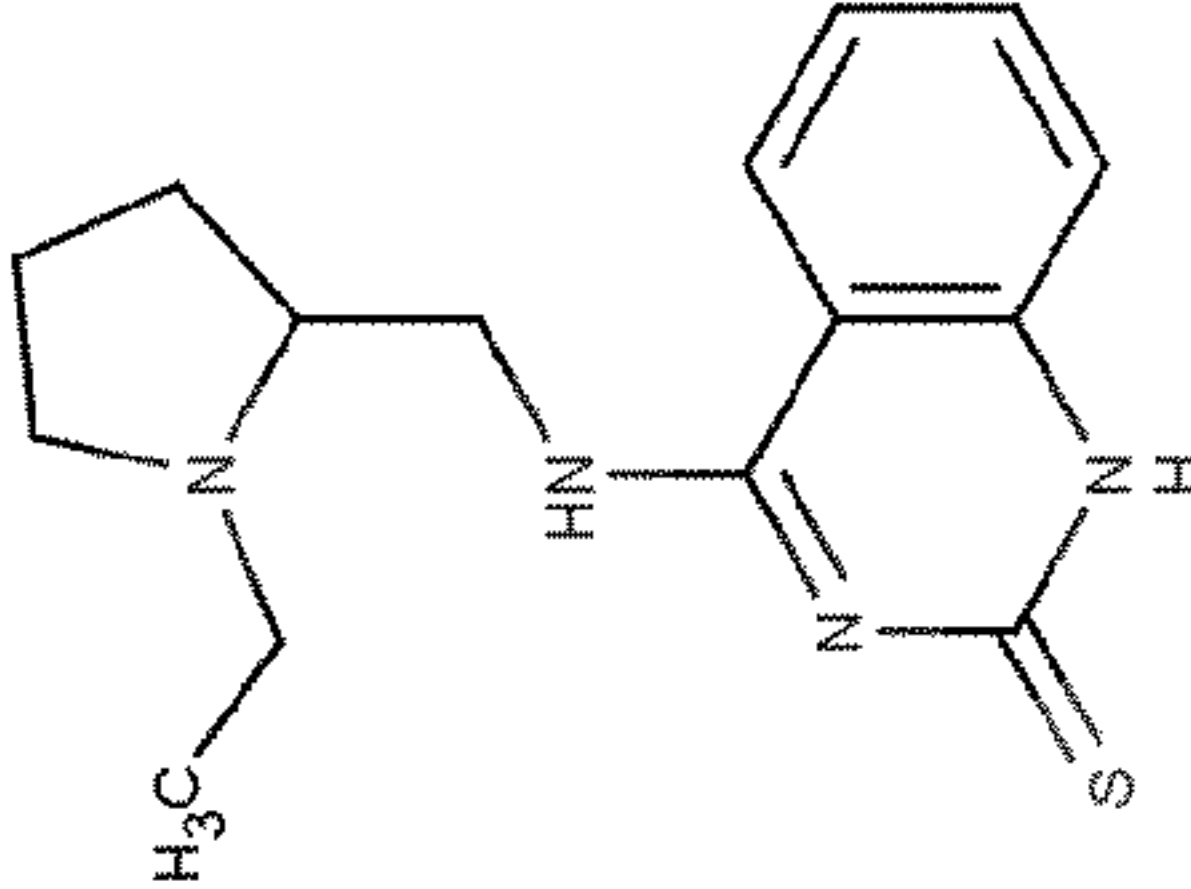
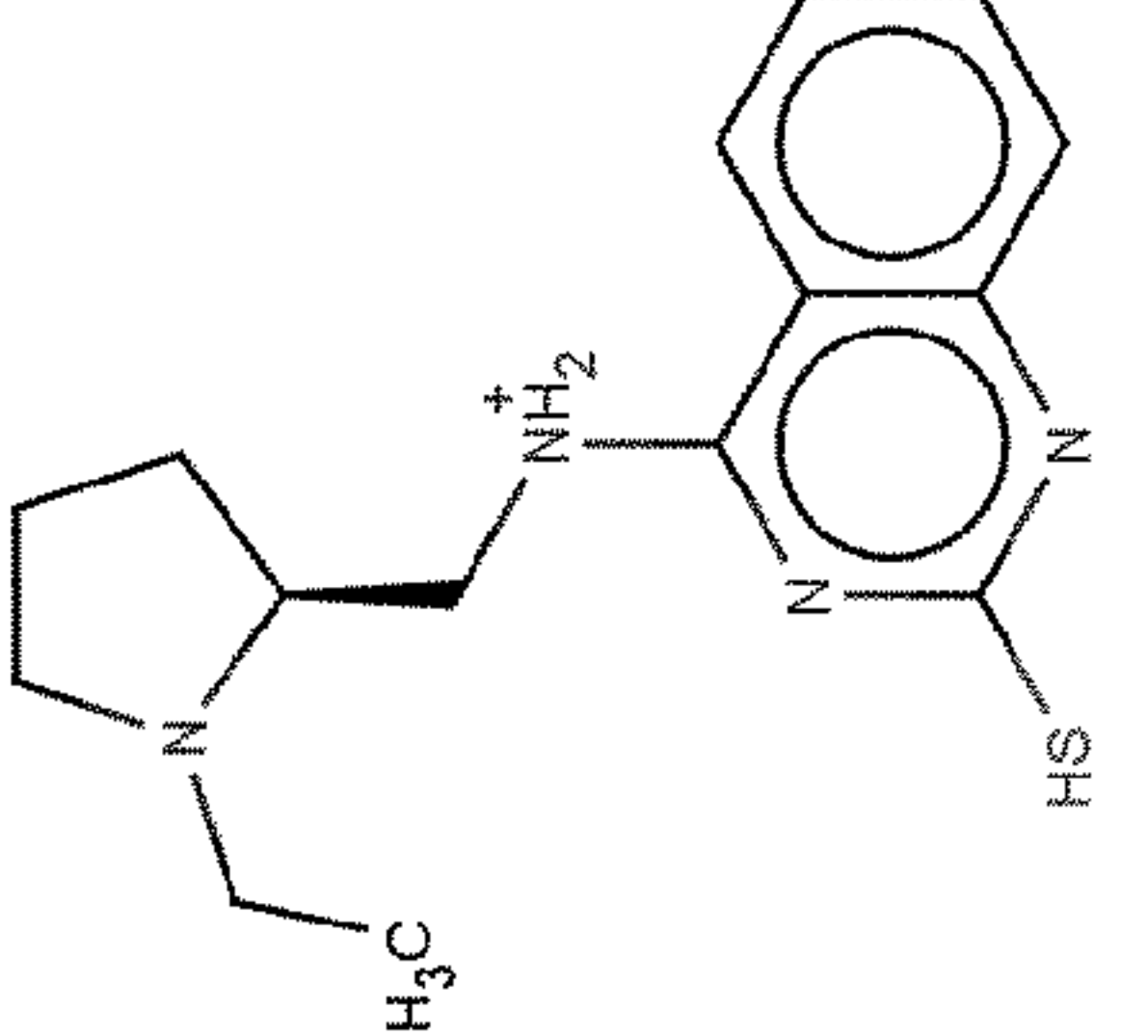
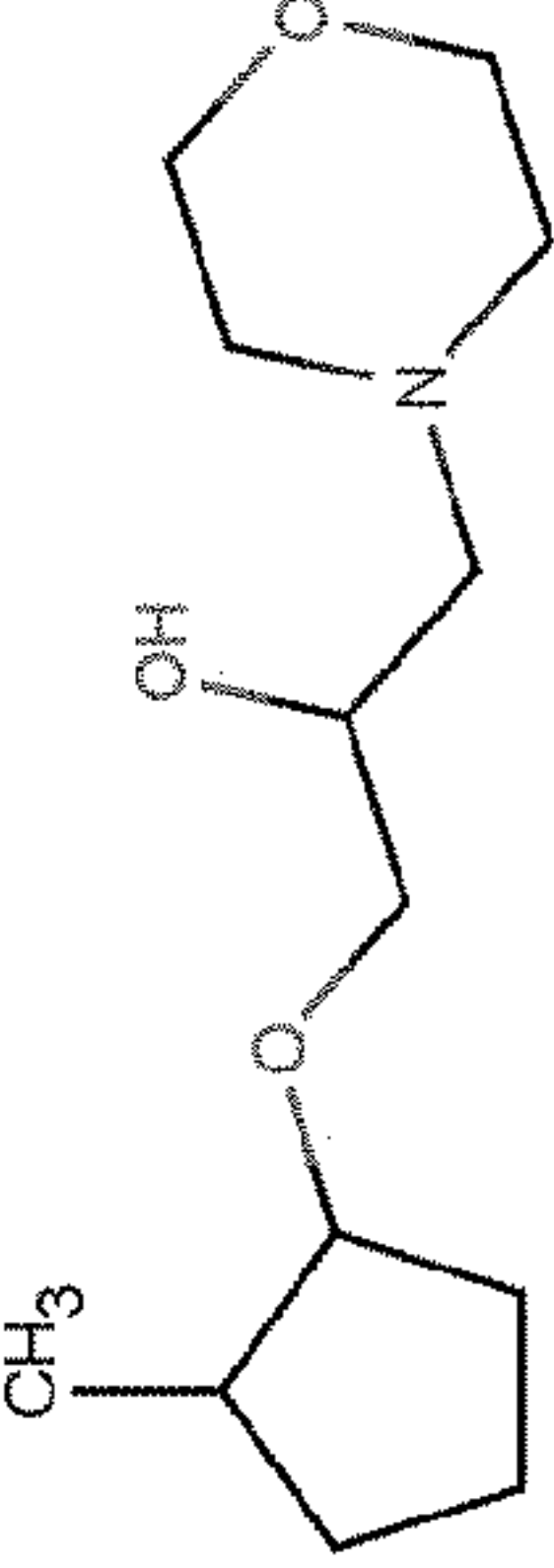
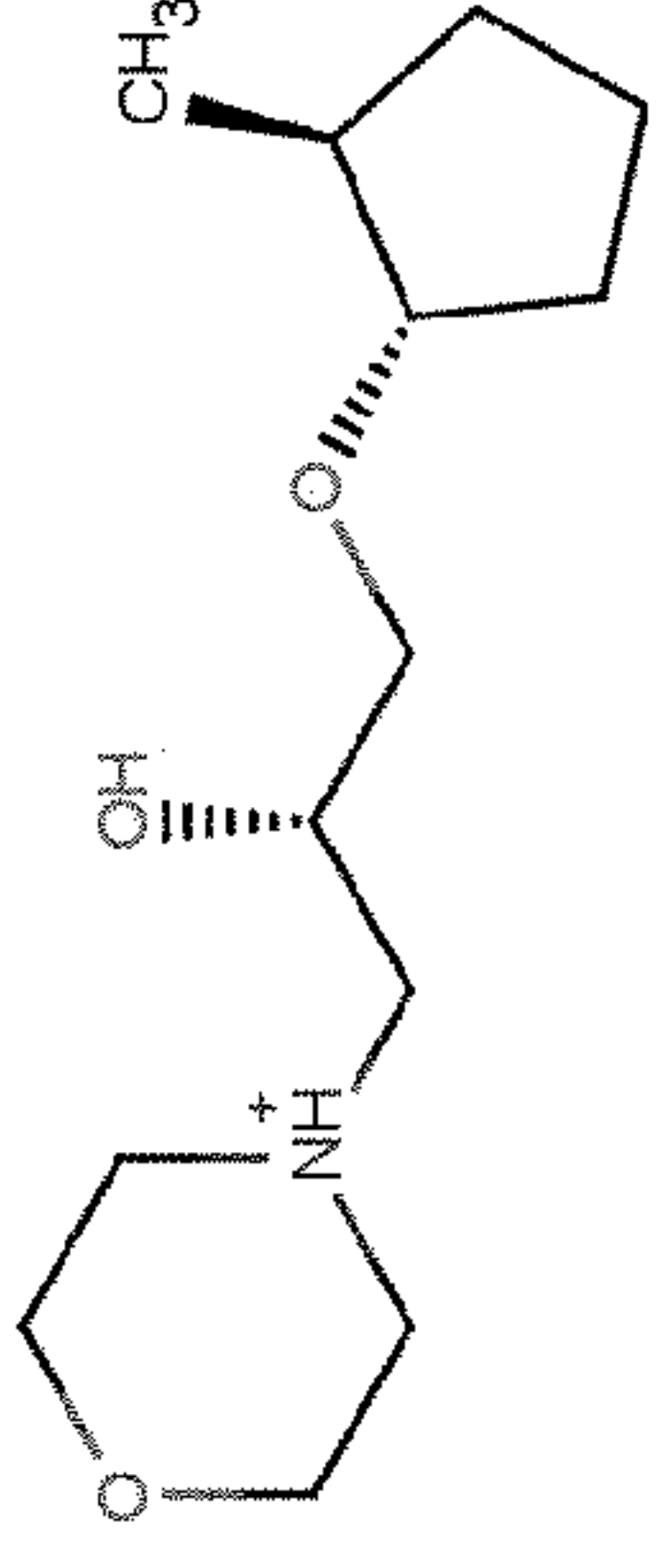
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 142		F2716-0877		F2716-0877_6_1 (1 of 8)
Compound 143		F2721-0110		F2721-0110_6_2 (1 of 14)
Compound 144		F2811-0081		F2811-0081_150_4 (1 of 451)

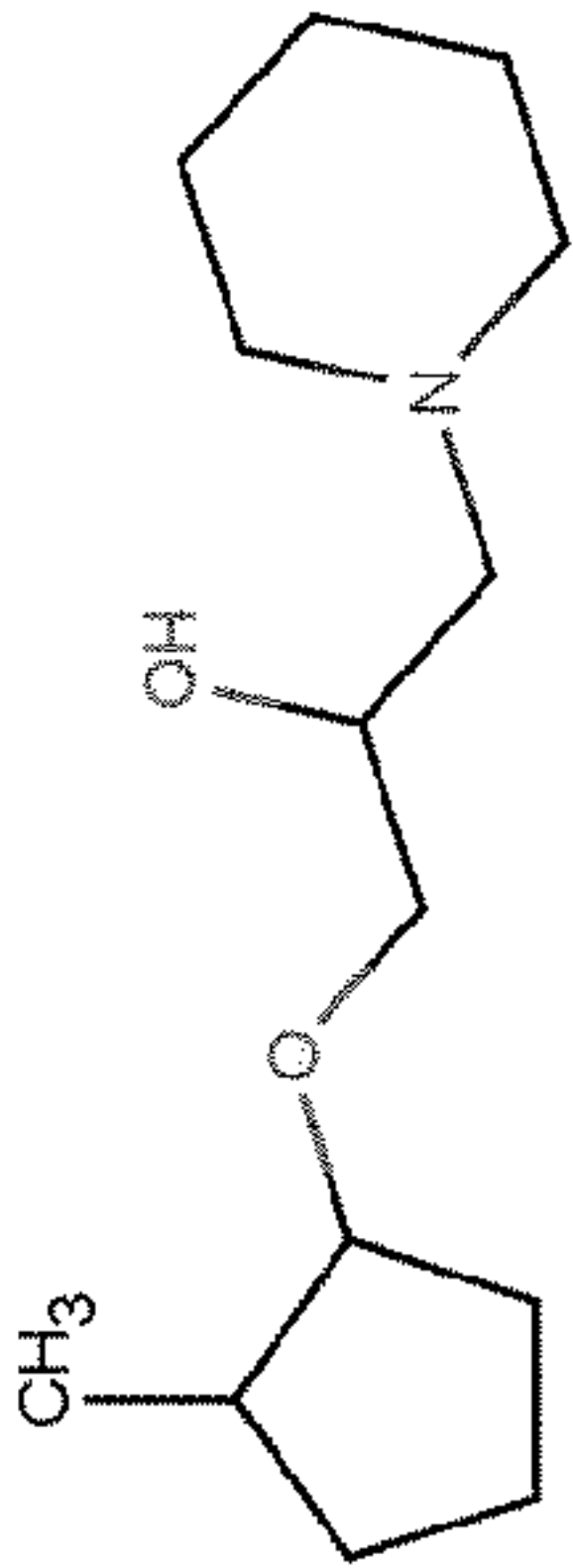
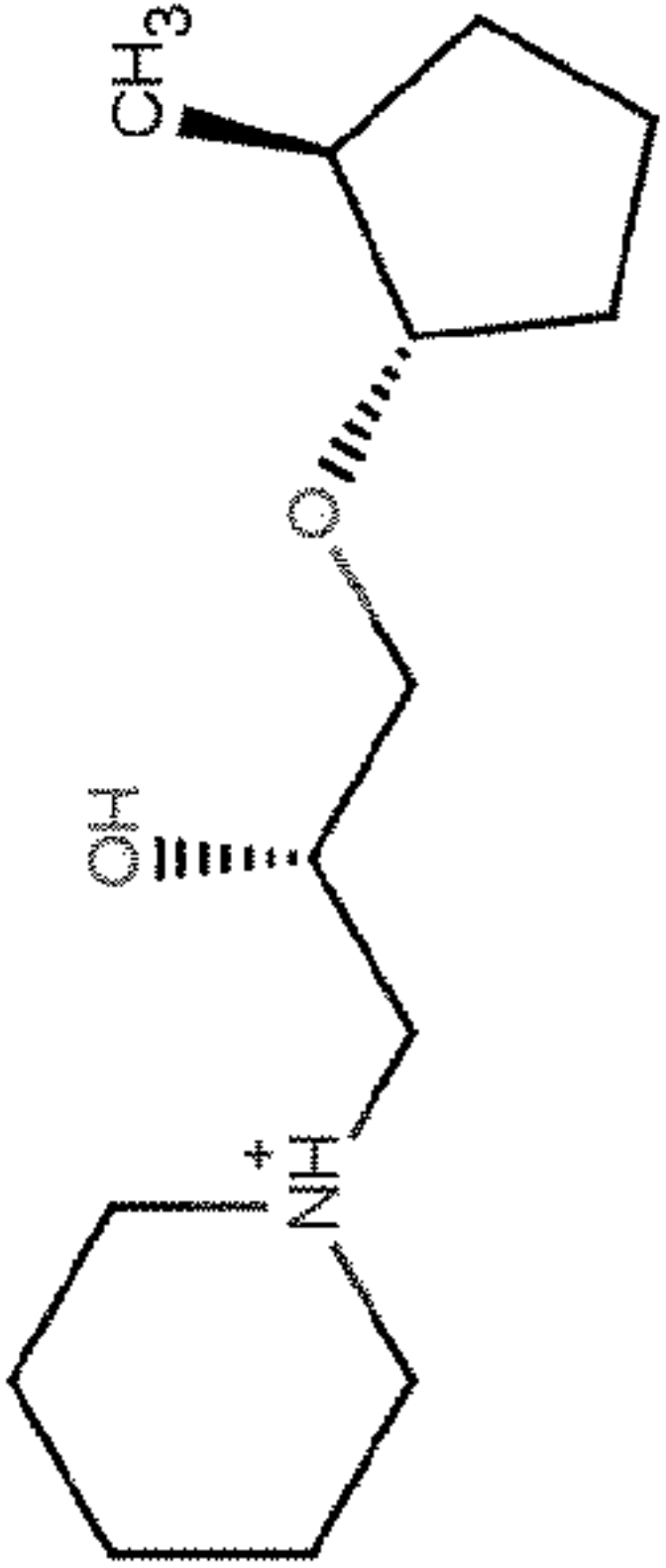
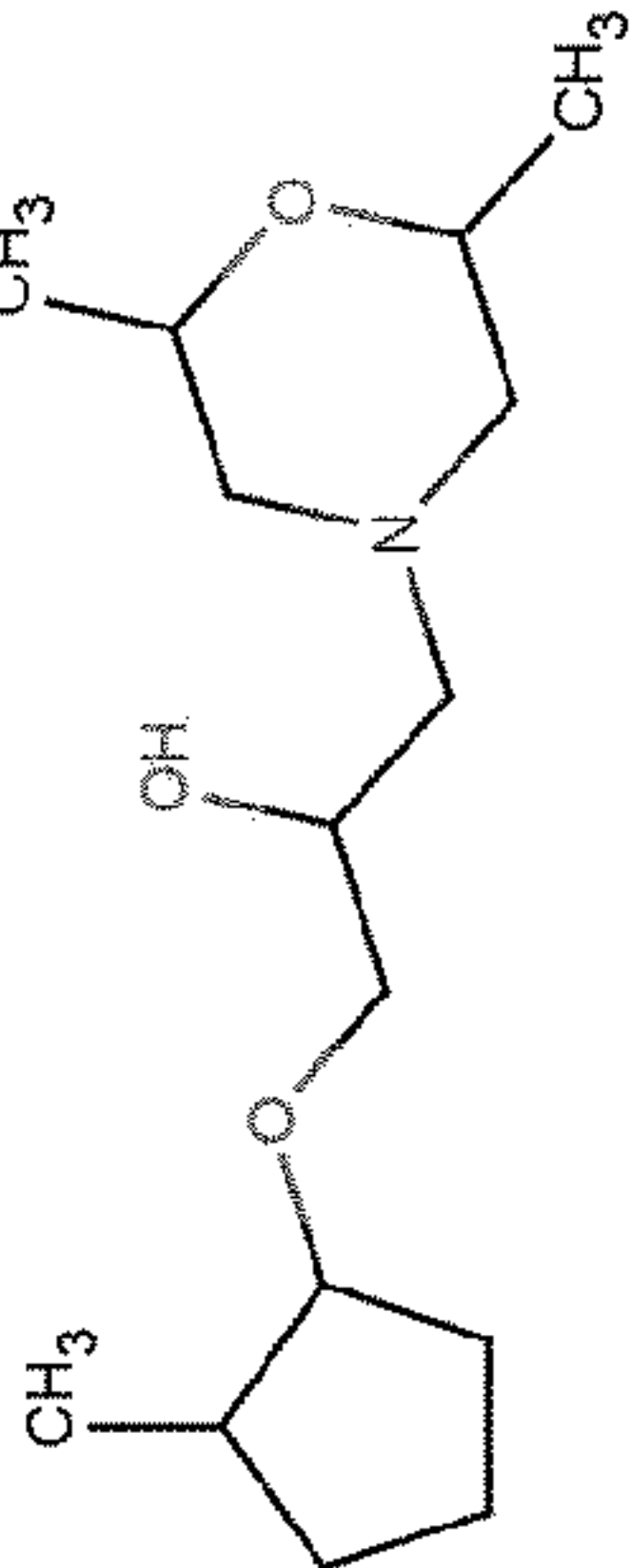
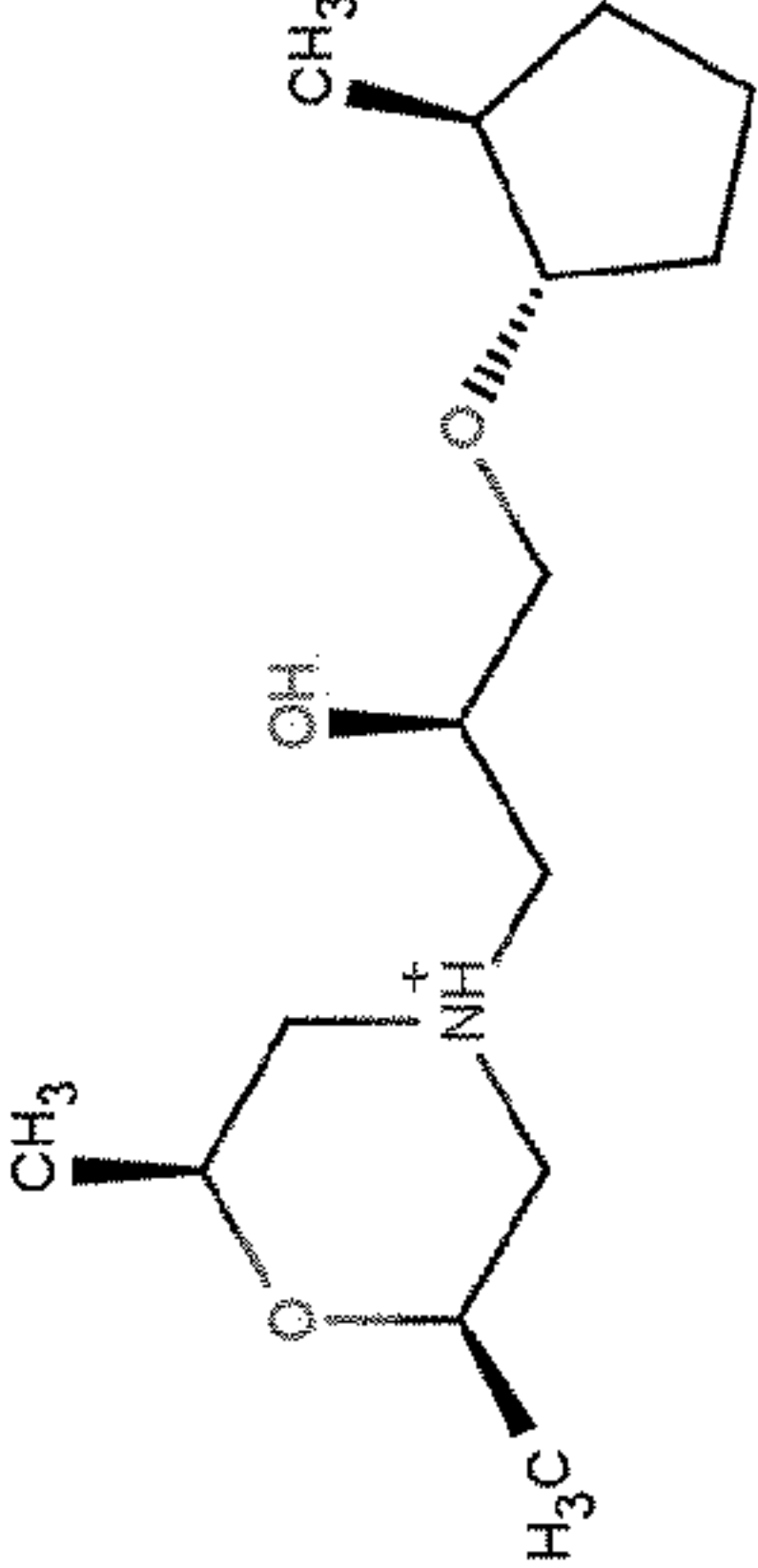
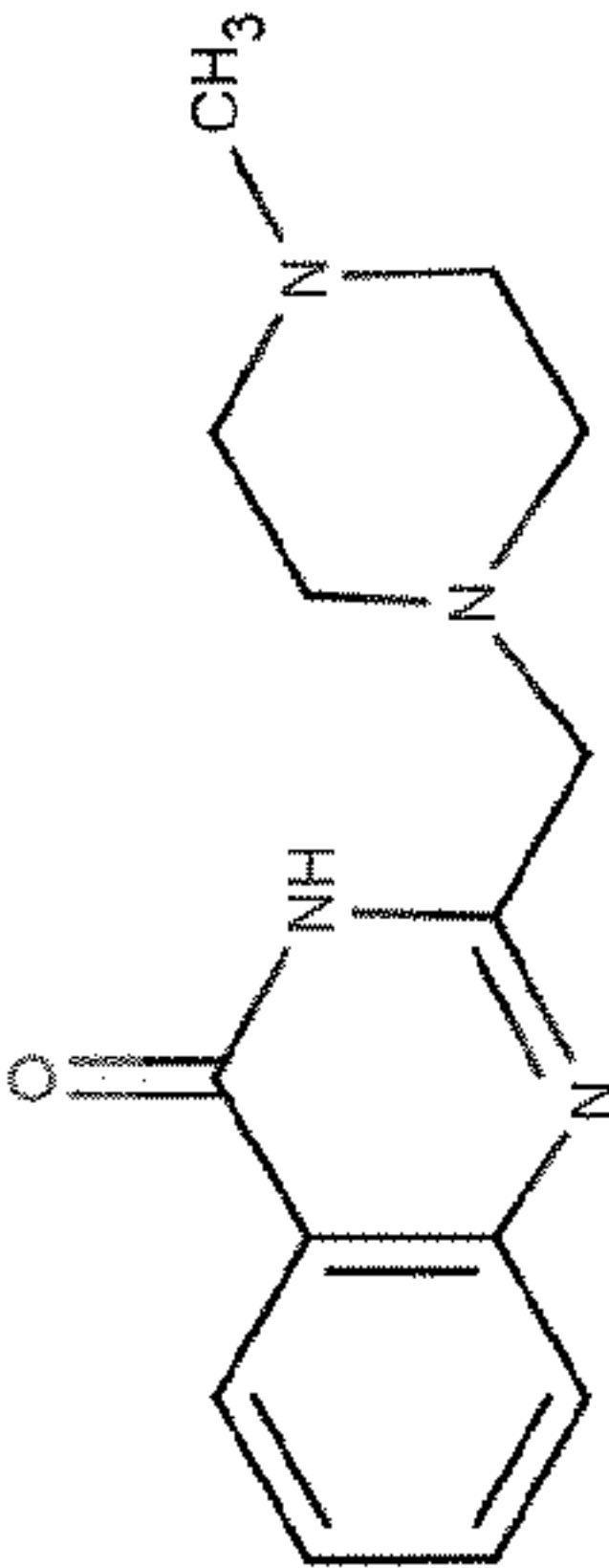
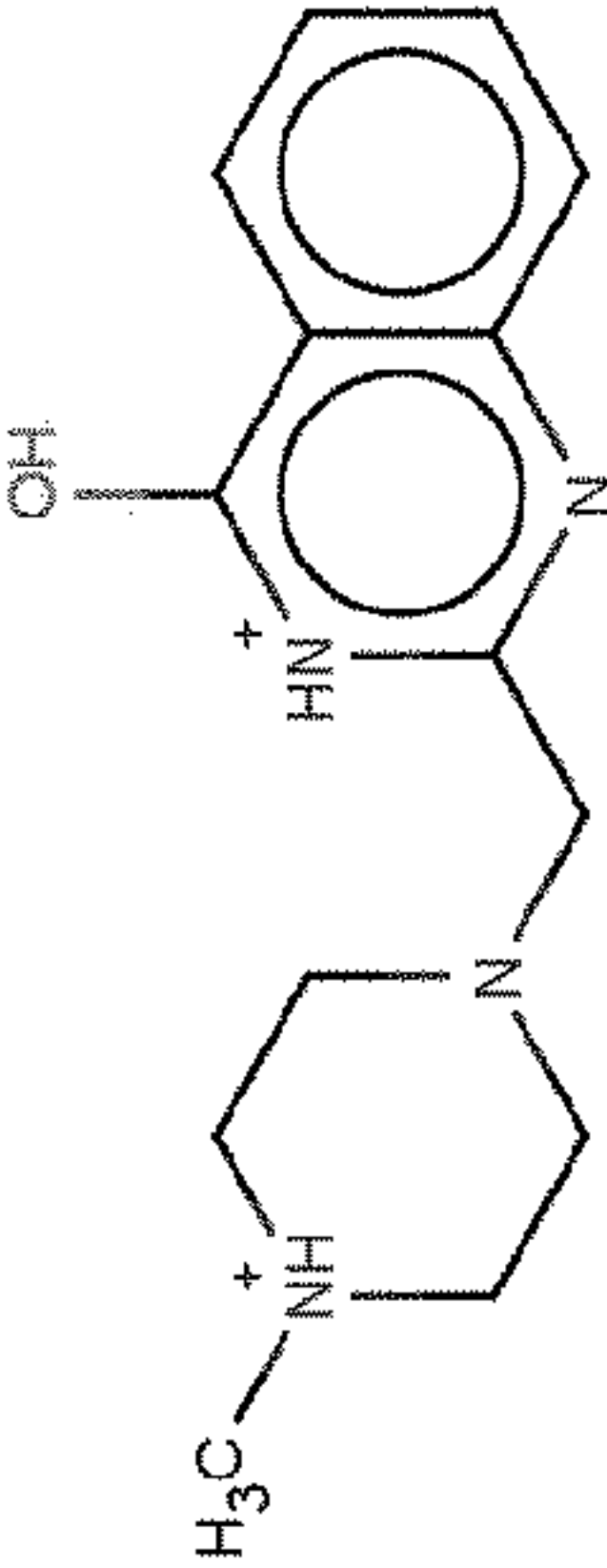
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 145		F2882-0062		F2882-0062_1_1 (1 of 21)
Compound 146		F3012-0022		F3012-0022_1_2 (1 of 4)
Compound 147		F3034-0039		F3034-0039_1 (1 of 8)

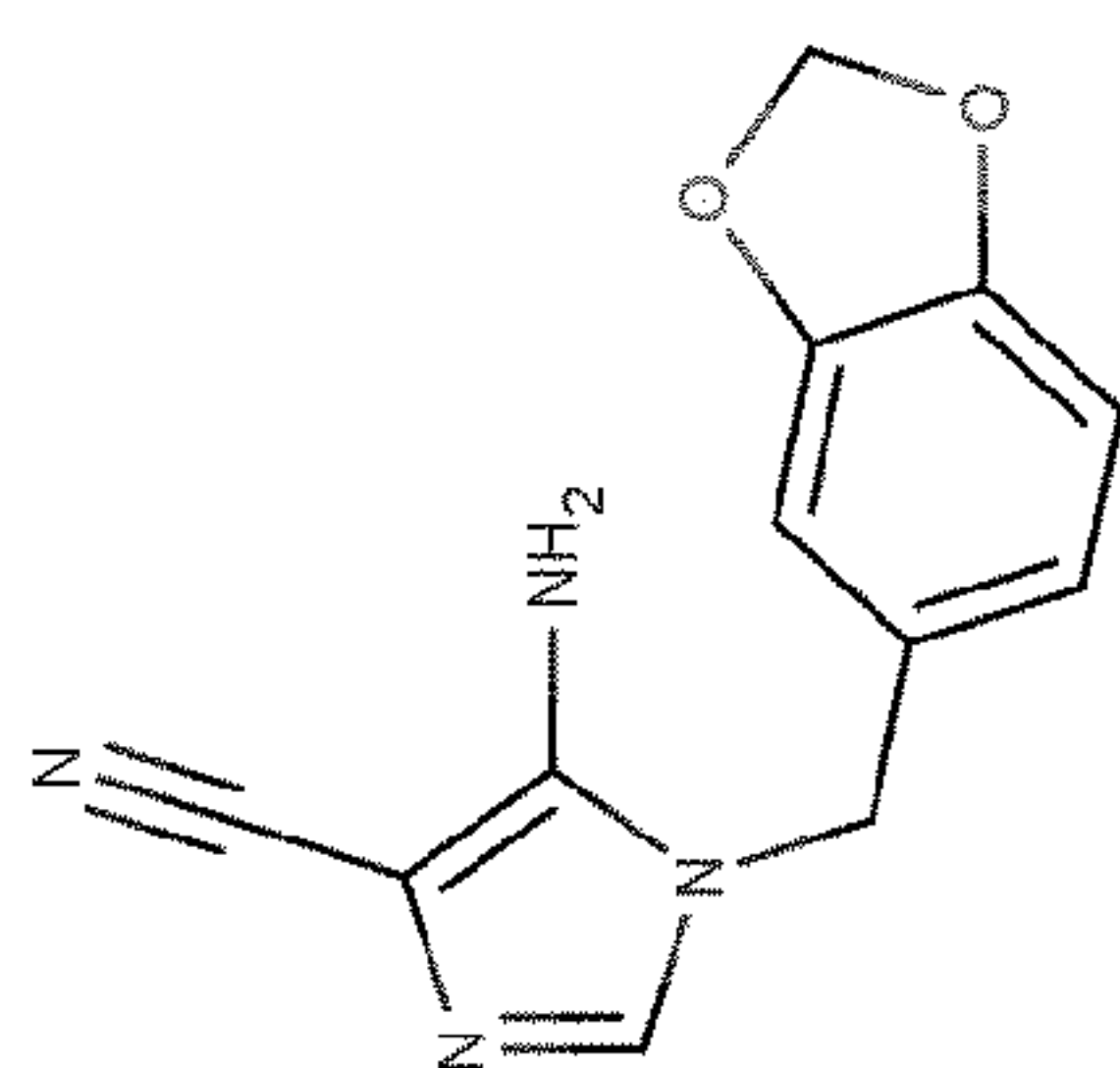
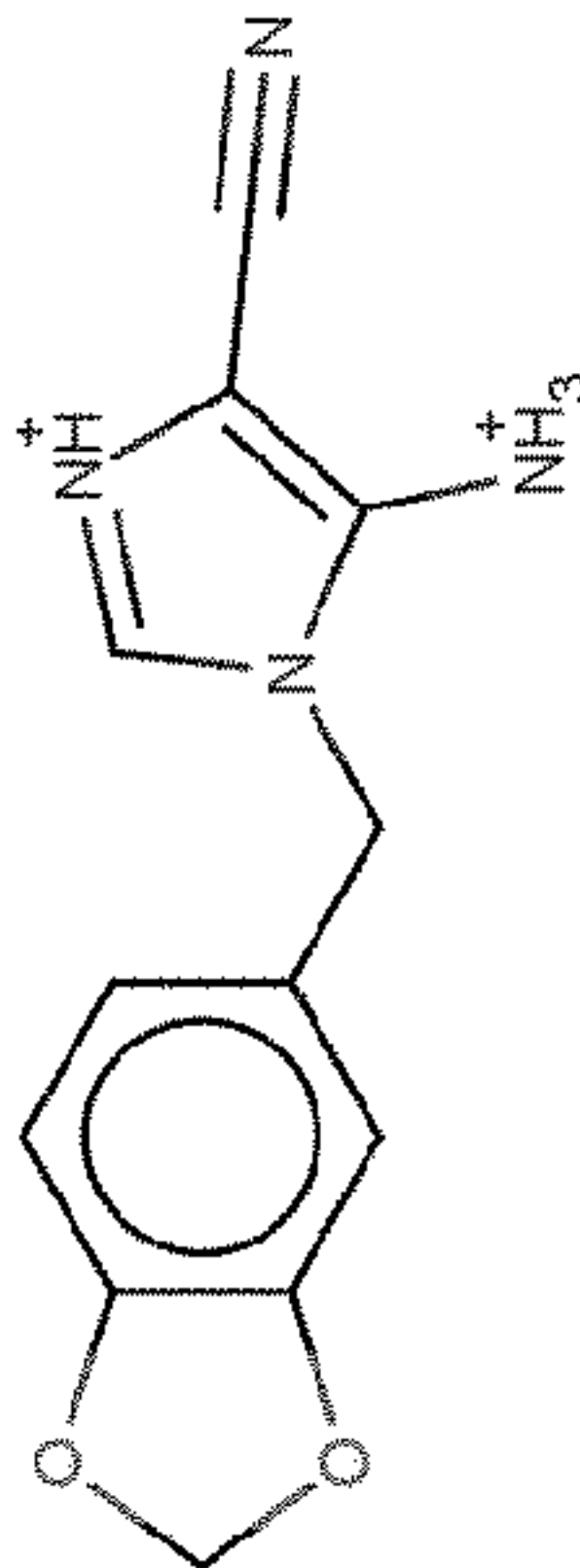
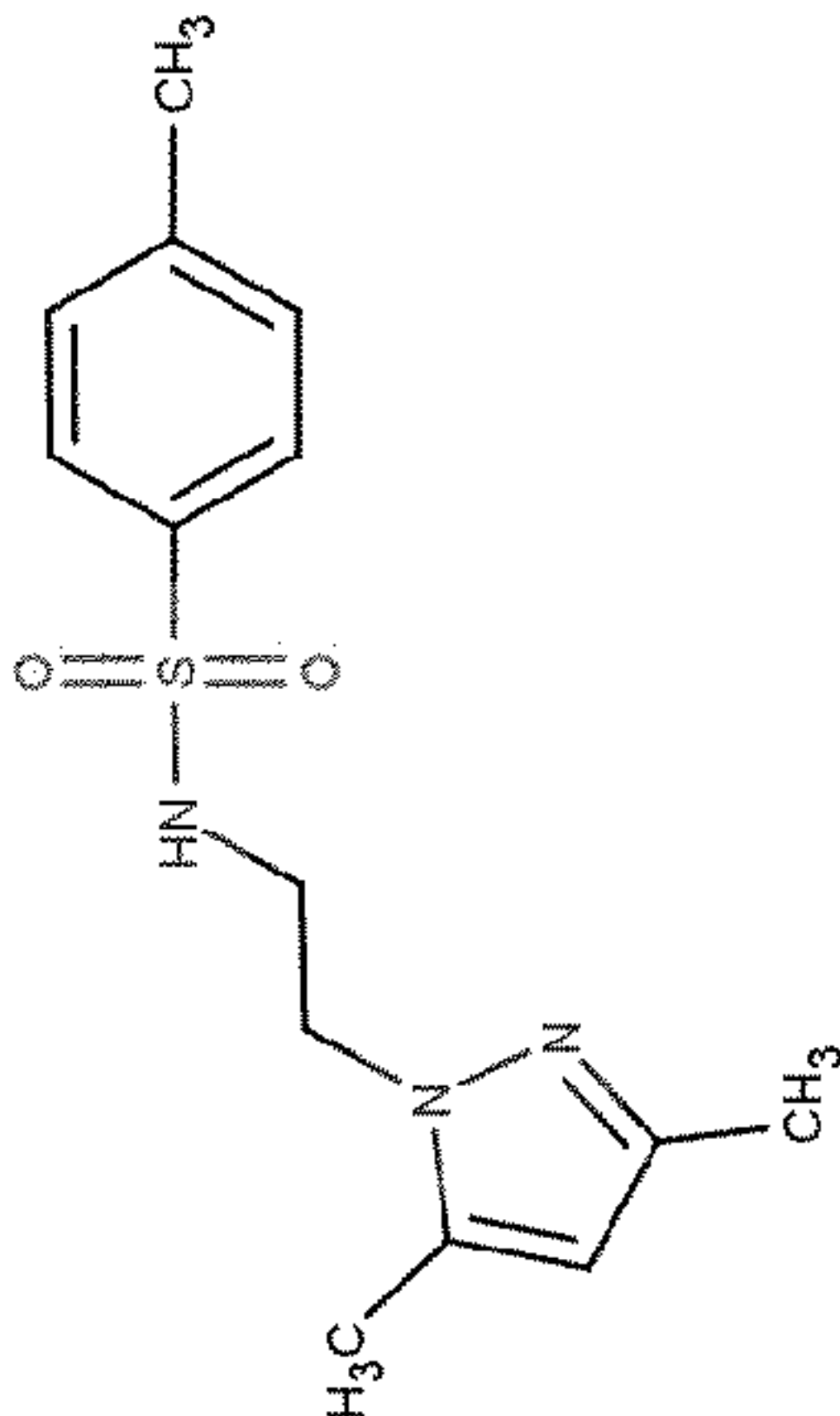
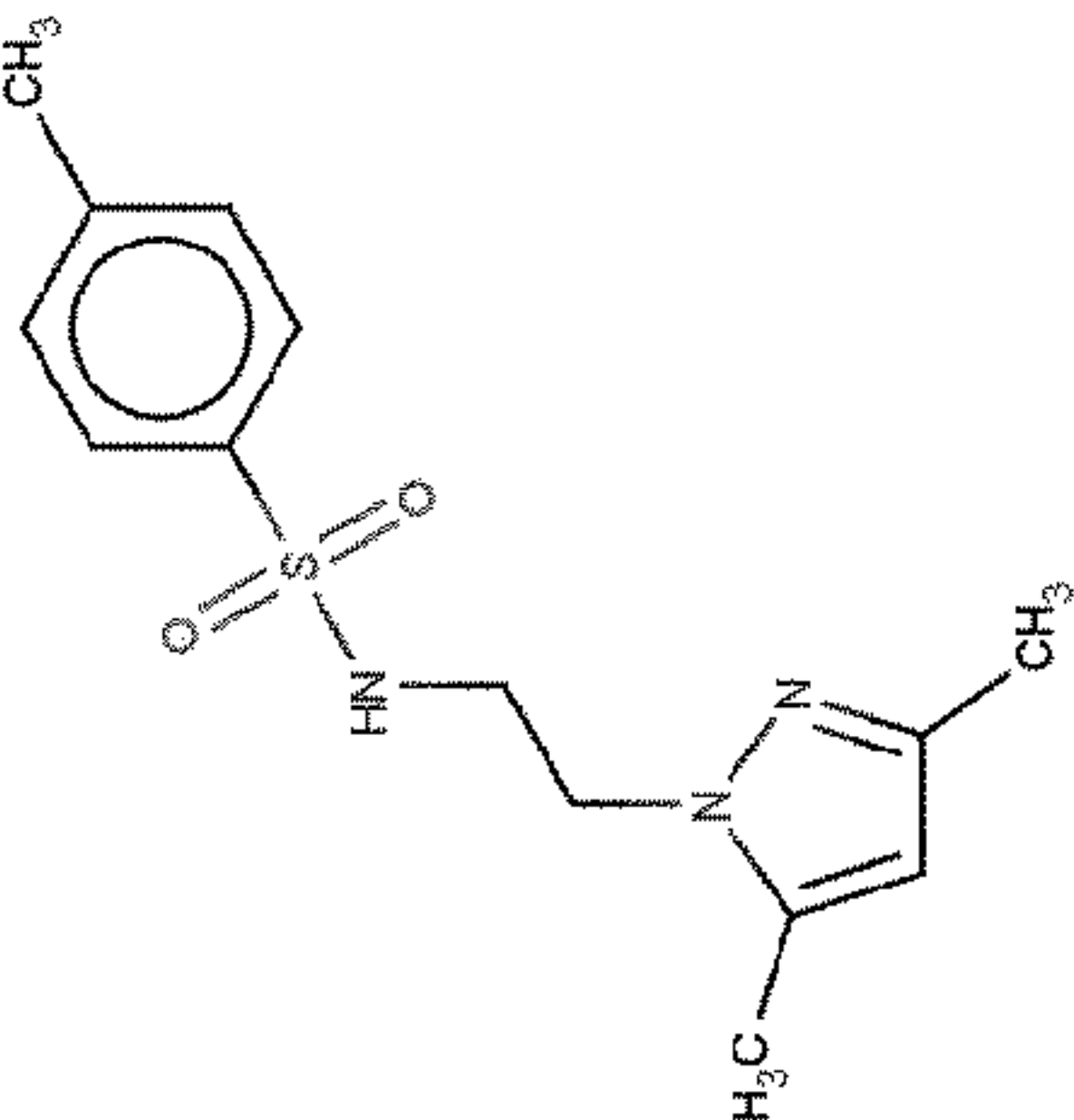
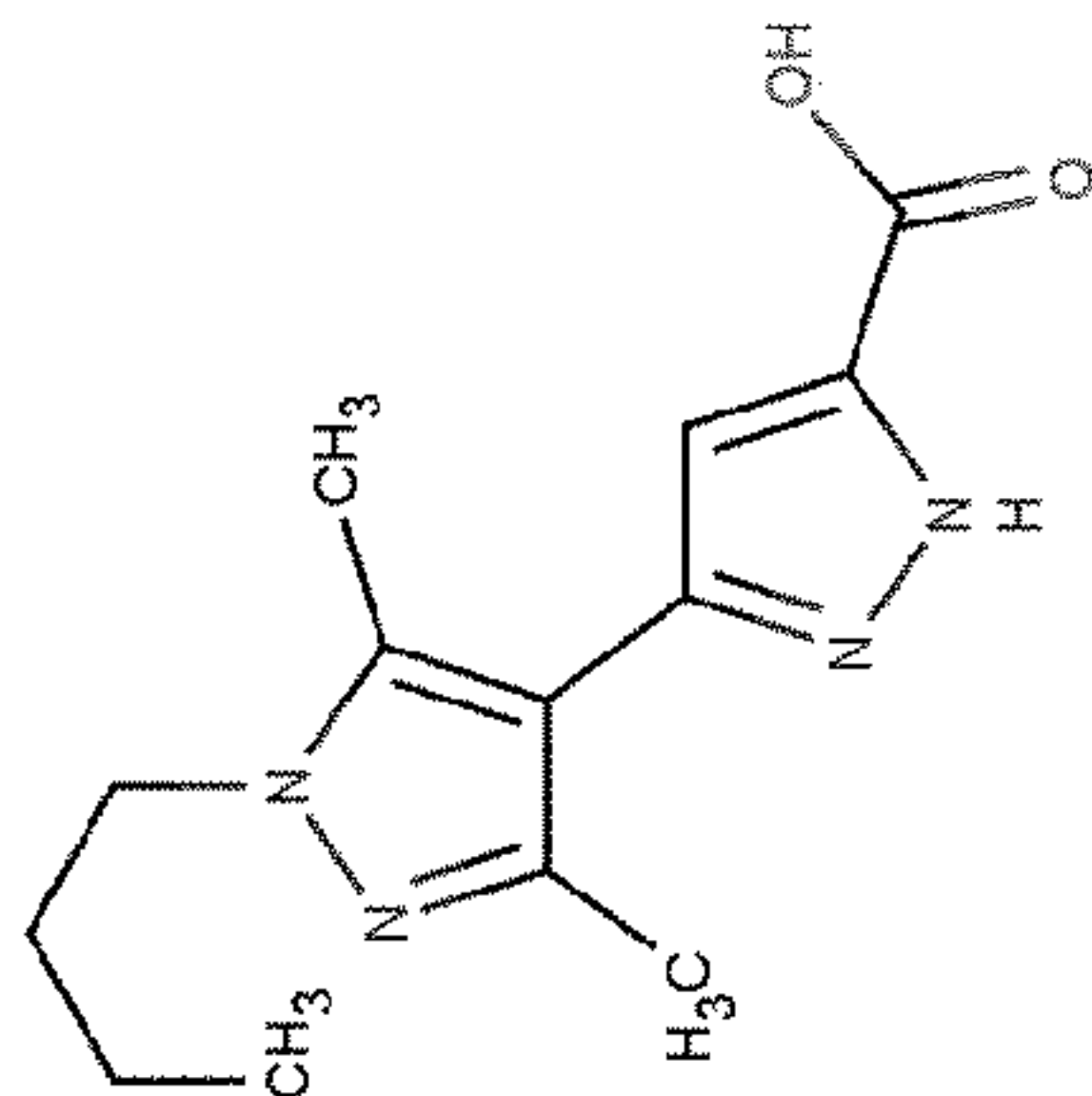
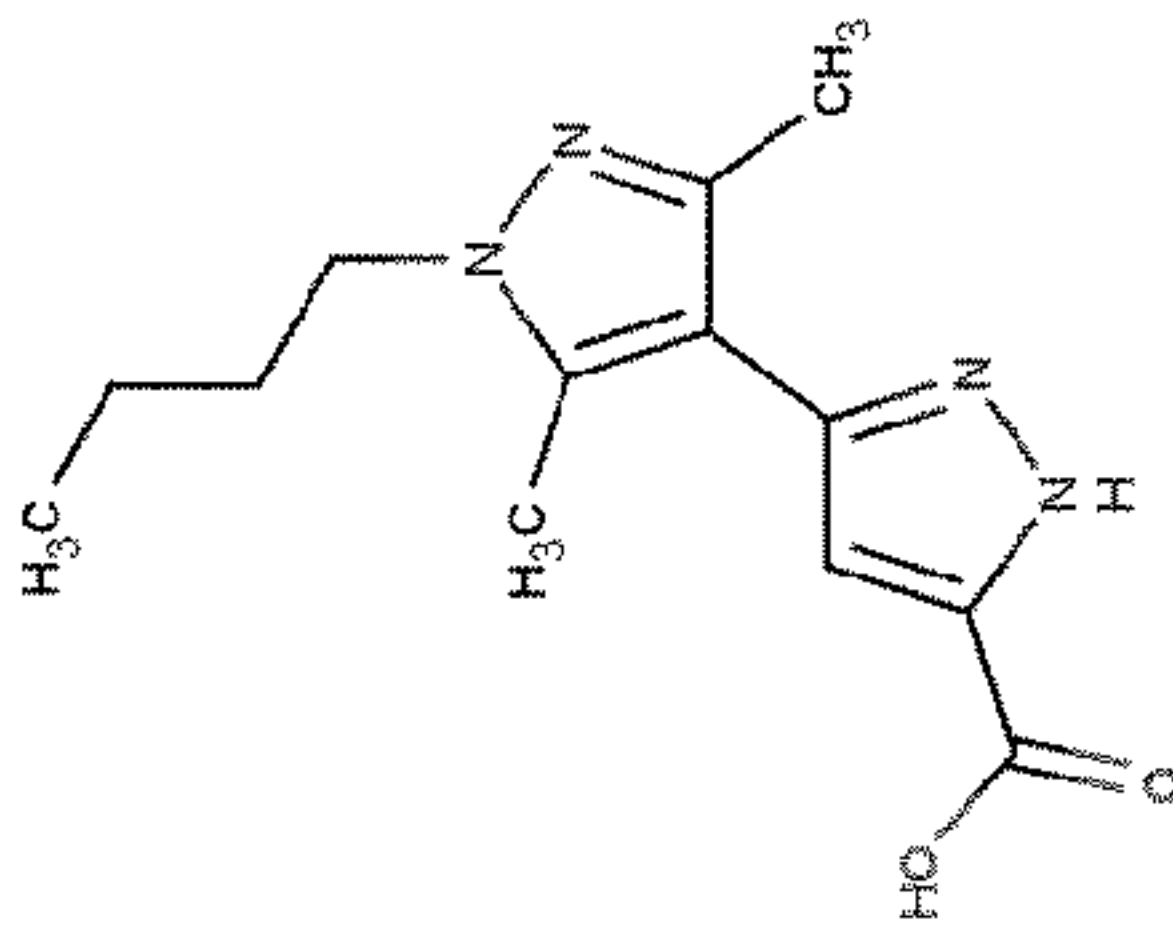
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 148		F3034-0286		F3034-0286_3_2 (1 of 4)
Compound 149		F3045-0001		F3045-0001_11_1 (1 of 14)
Compound 150		F3084-0058		F3084-0058_2_1 (1 of 4)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 151		F3084-0157		F3084-0157_2_2 (1 of 8)
Compound 152		F3099-0269		F3099-0269_1_1 (1 of 8)
Compound 153		F3099-3404		F3099-3404_29_2 (1 of 120)

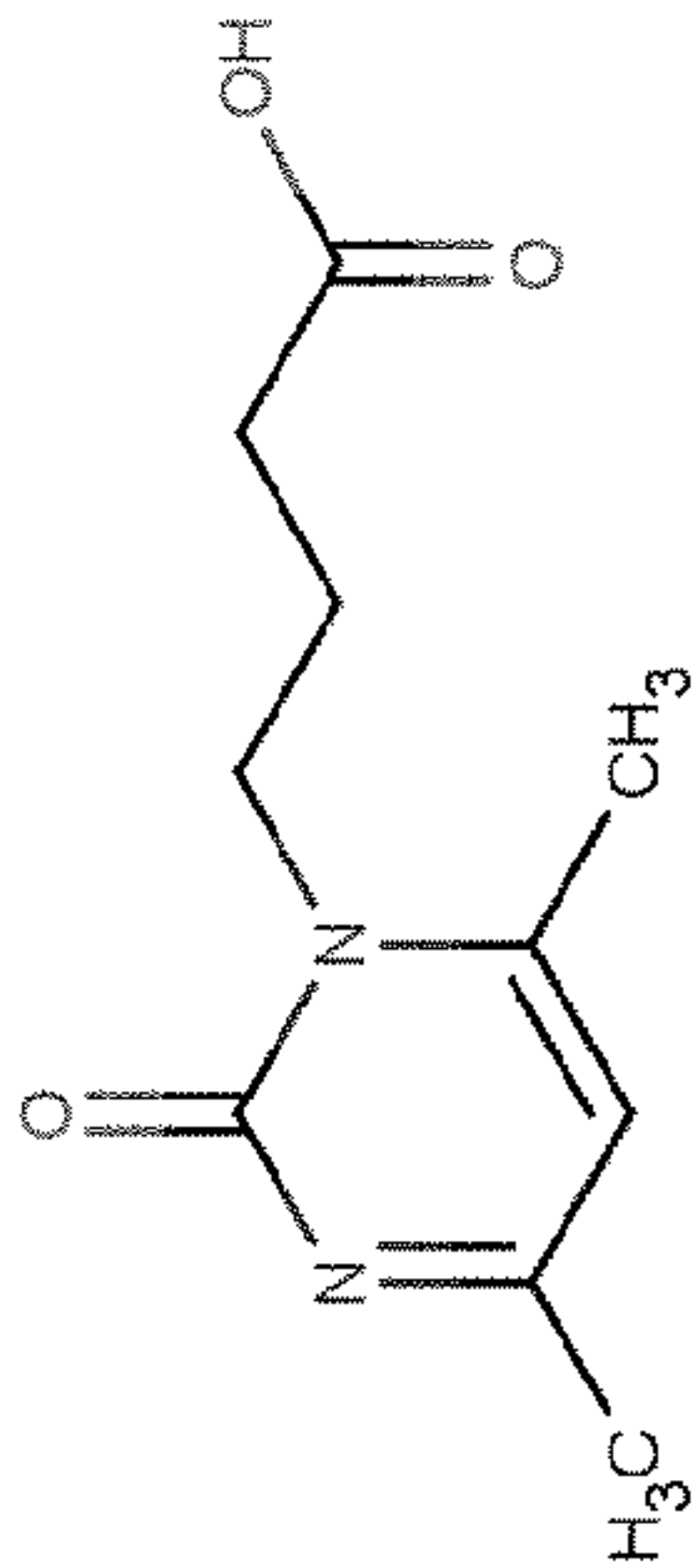
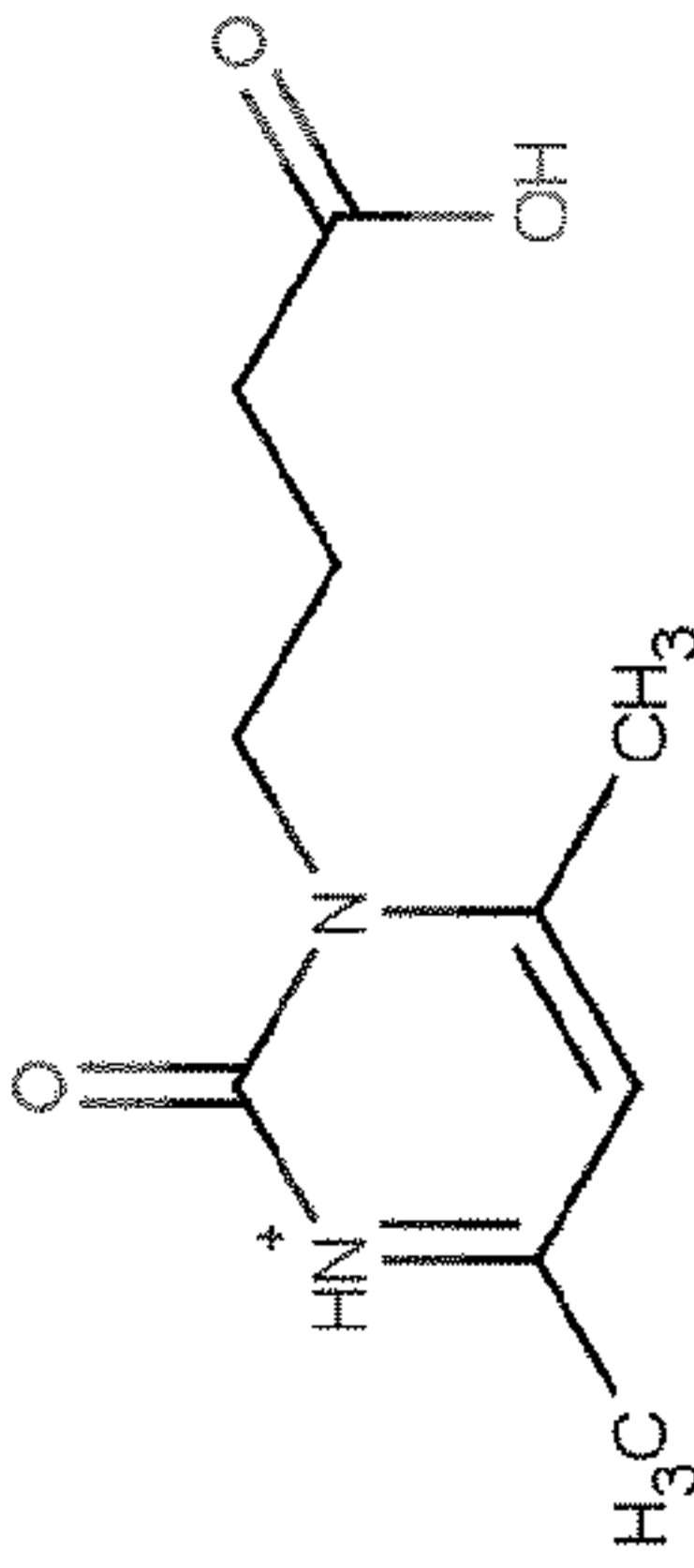
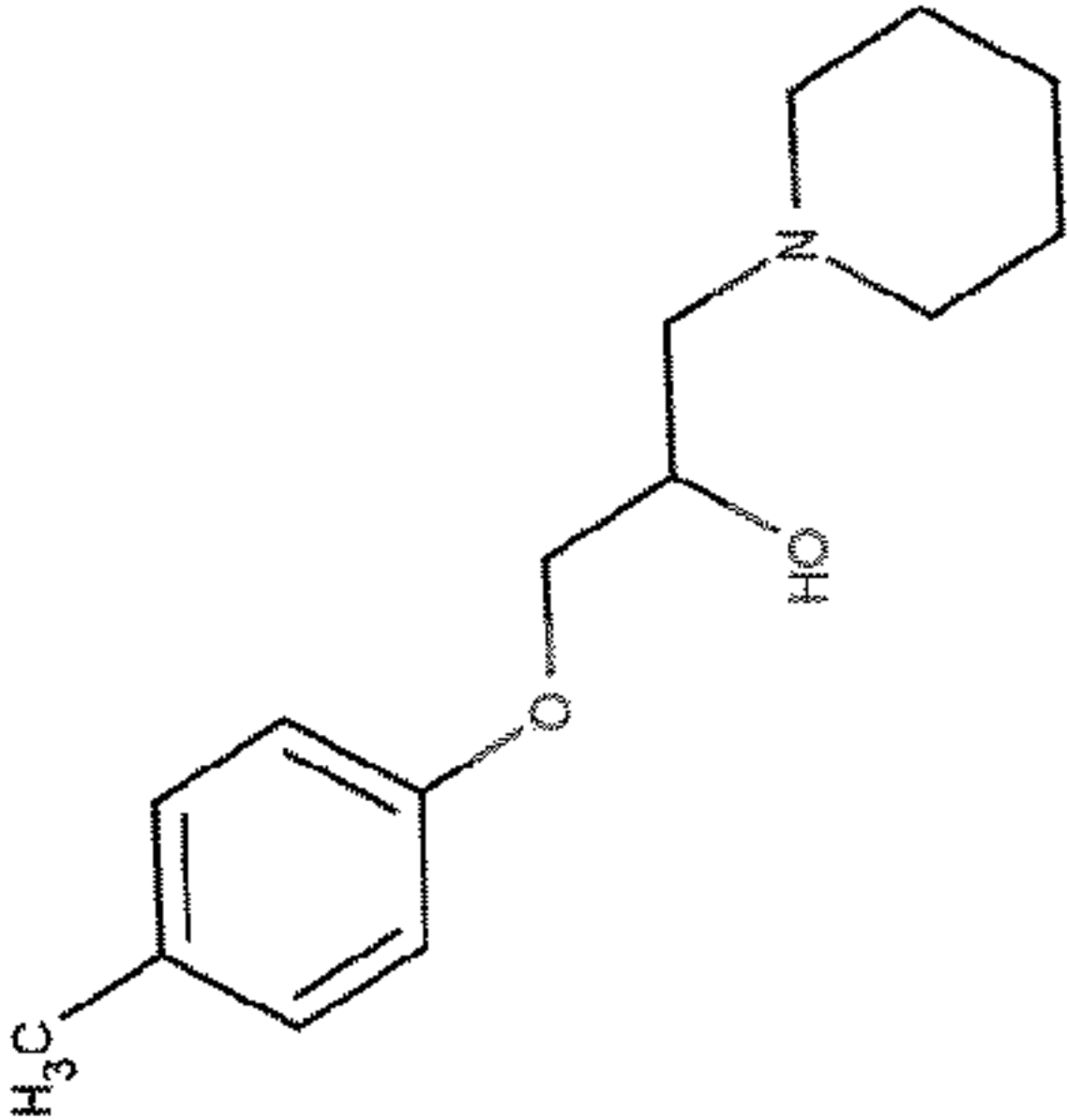
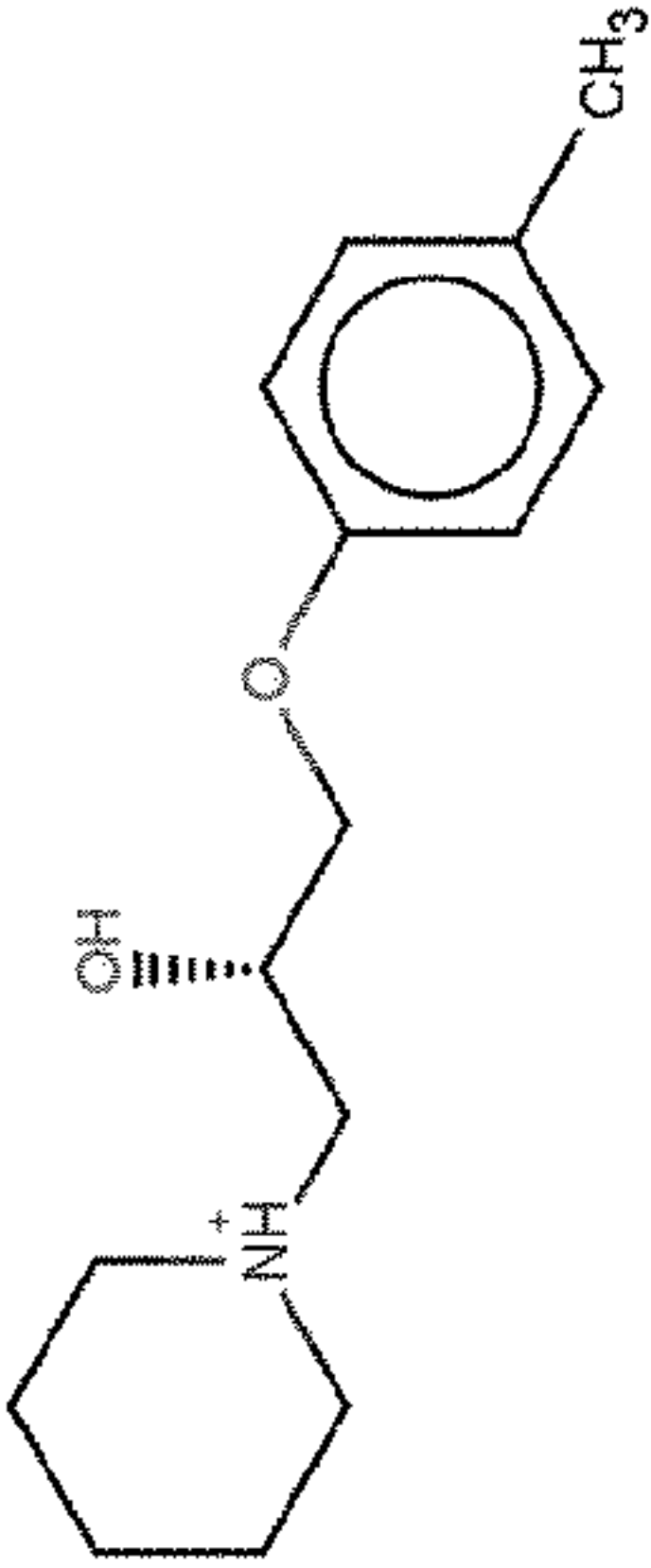
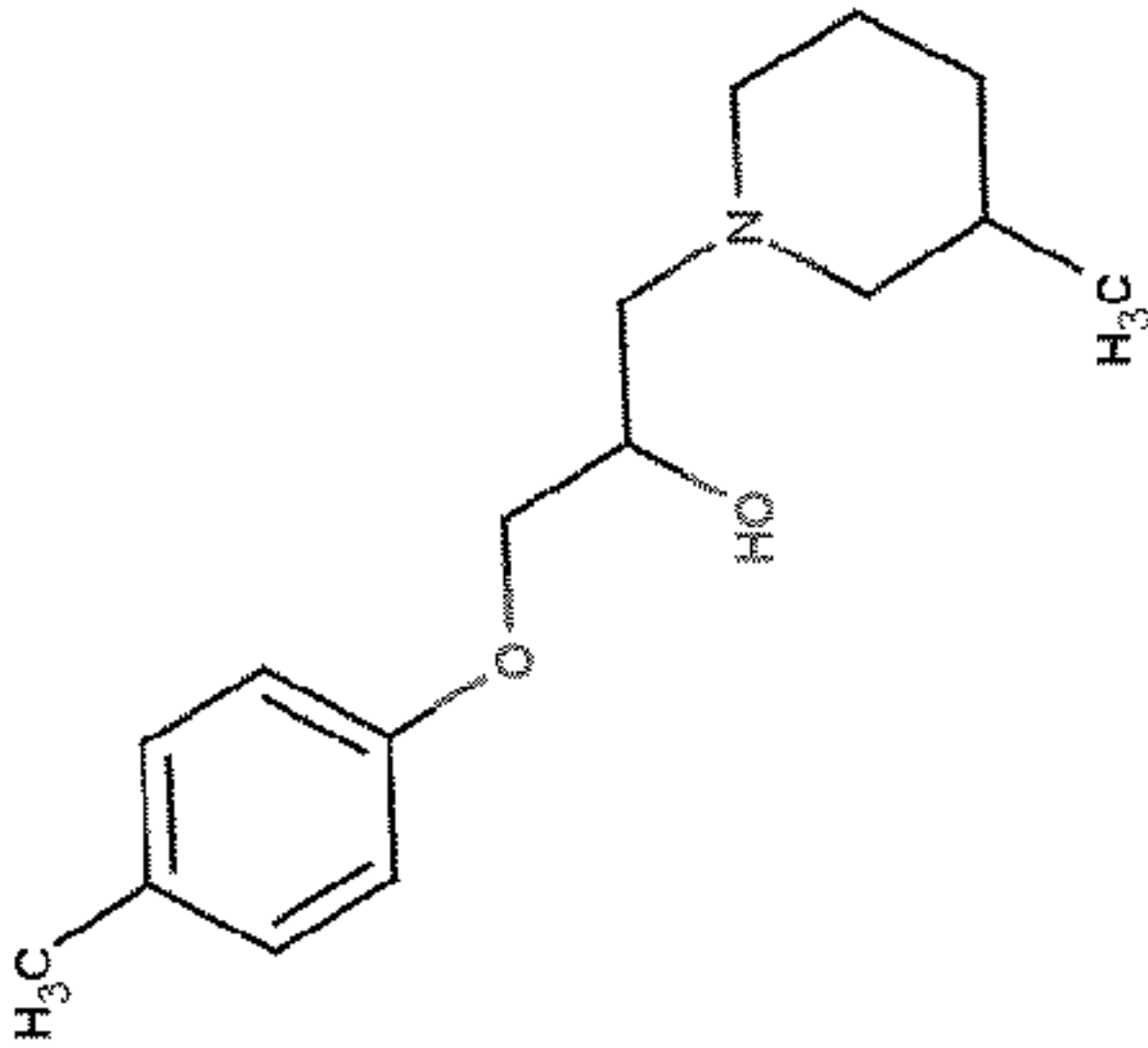
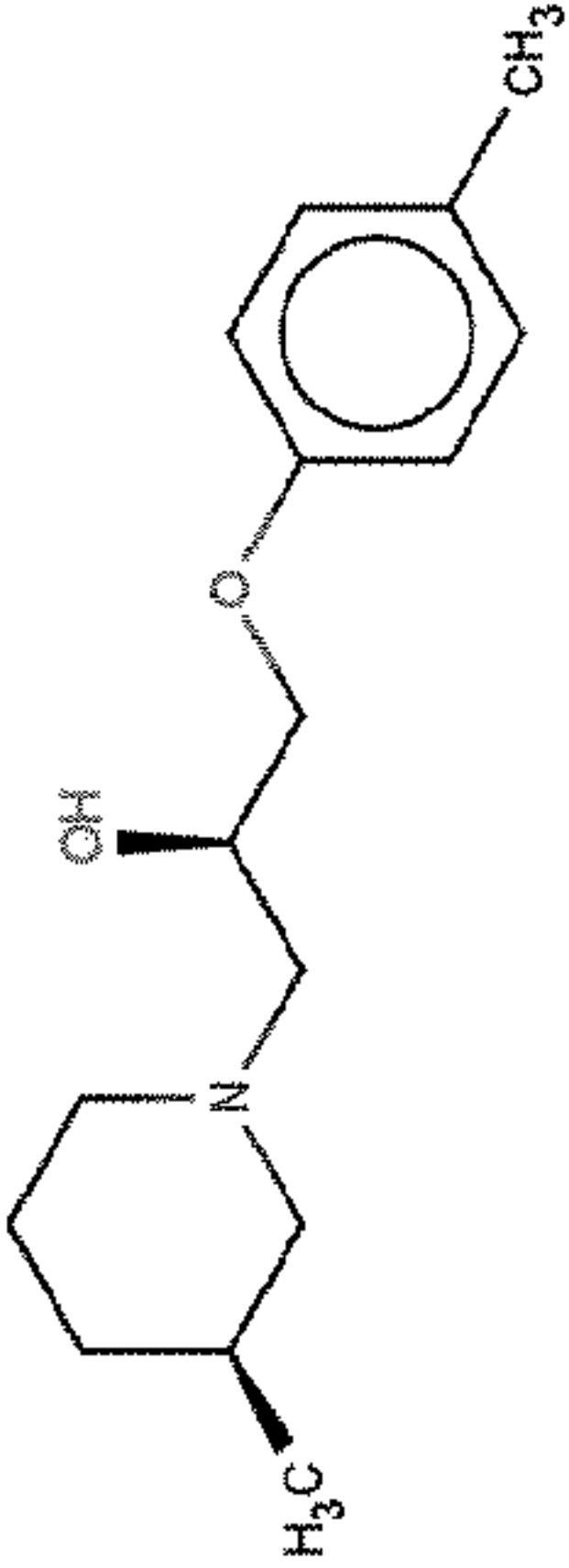
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 154		F3099-3503		F3099-3503_1_2 (1 of 24)
Compound 155		F3129-0147		F3129-0147_6_3 (1 of 24)
Compound 156		F3146-0231		F3146-0231_2_1 (1 of 8)

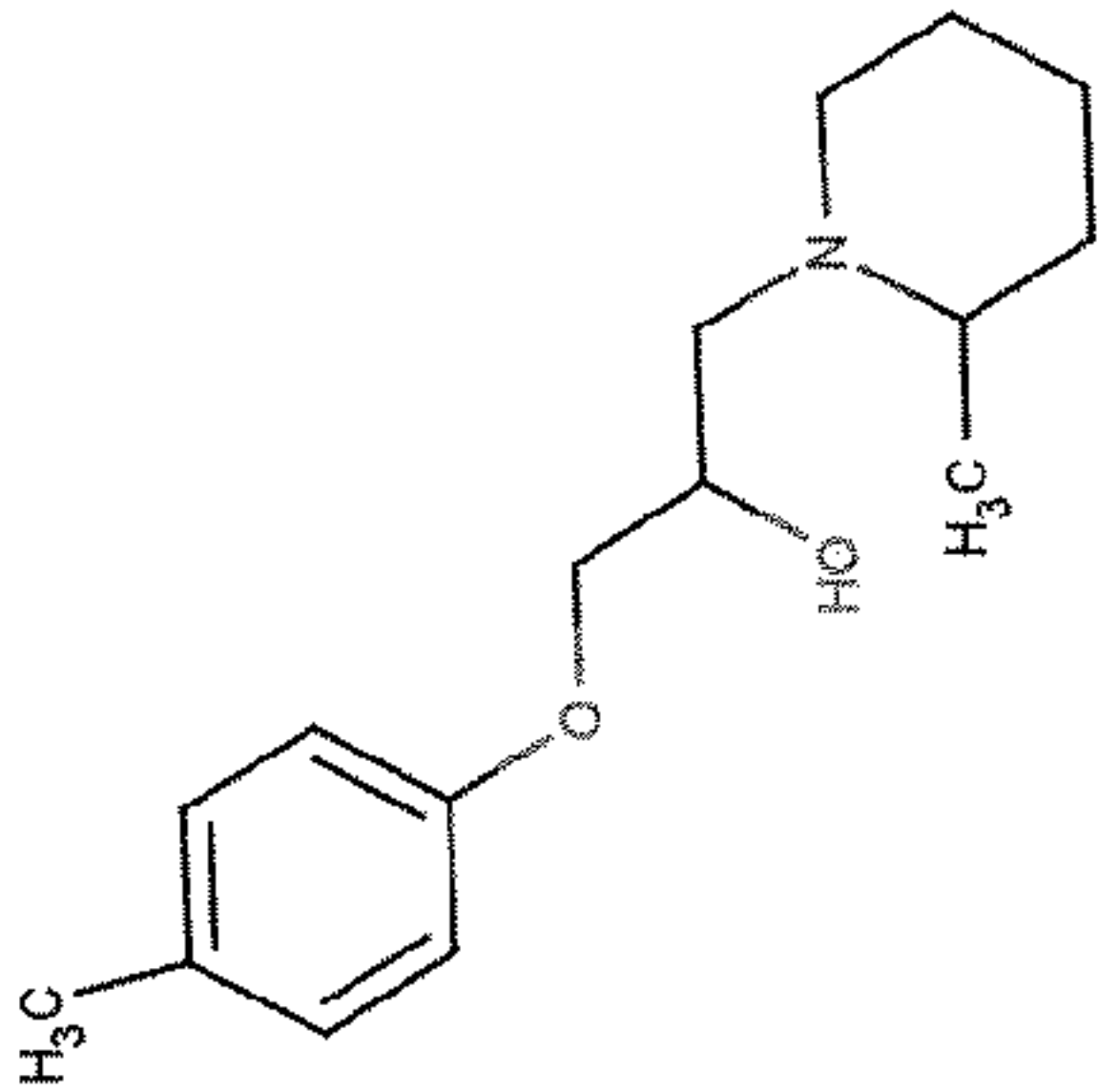
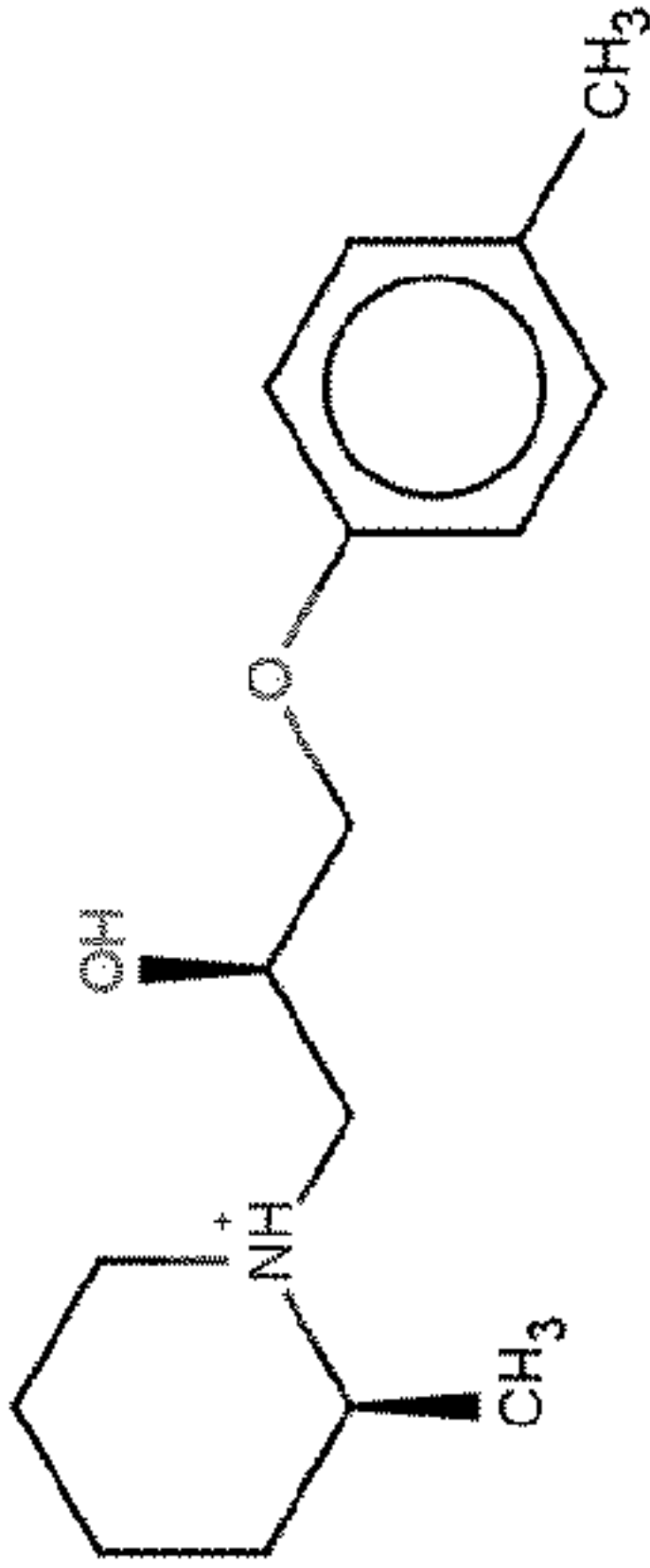
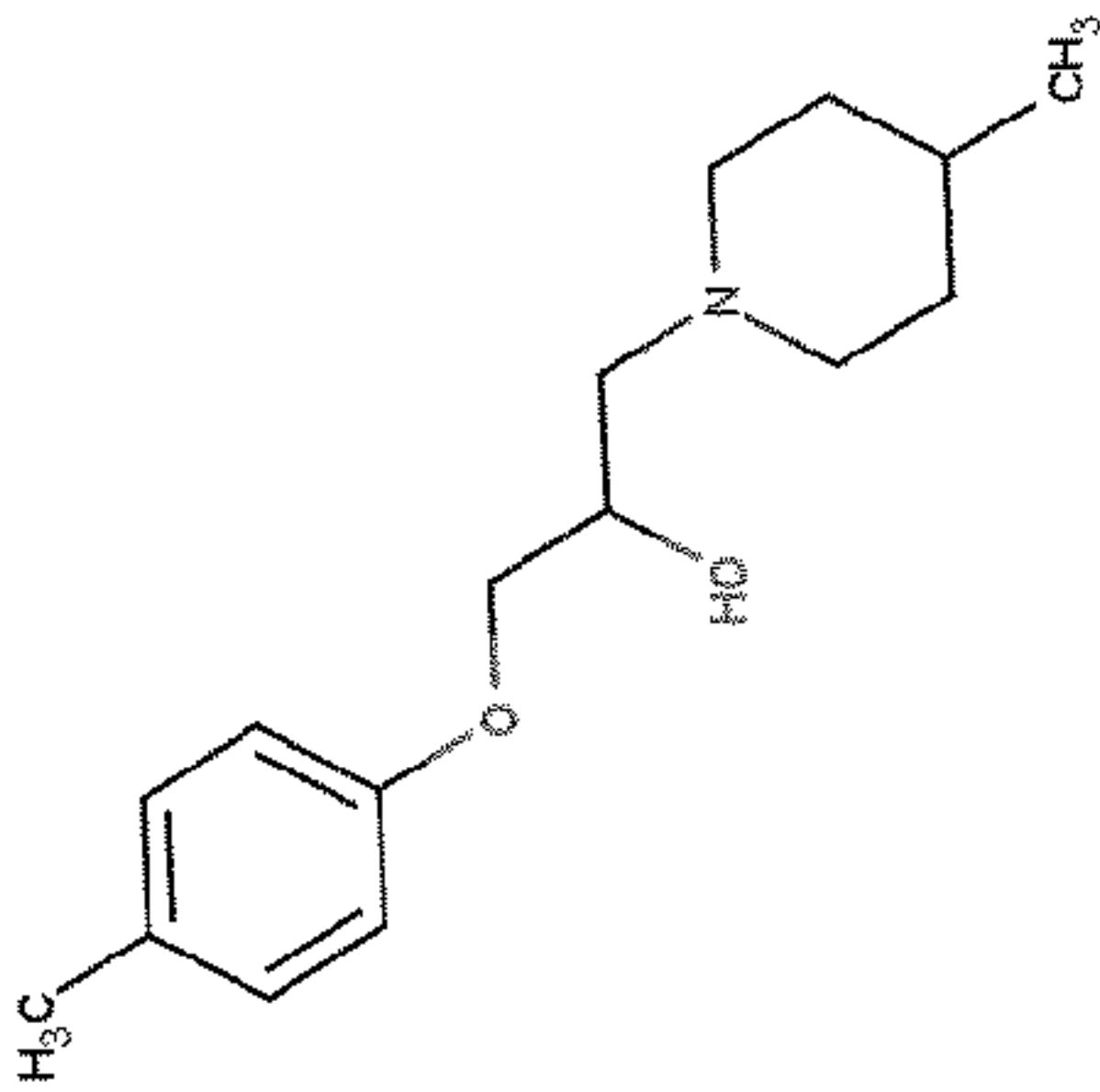
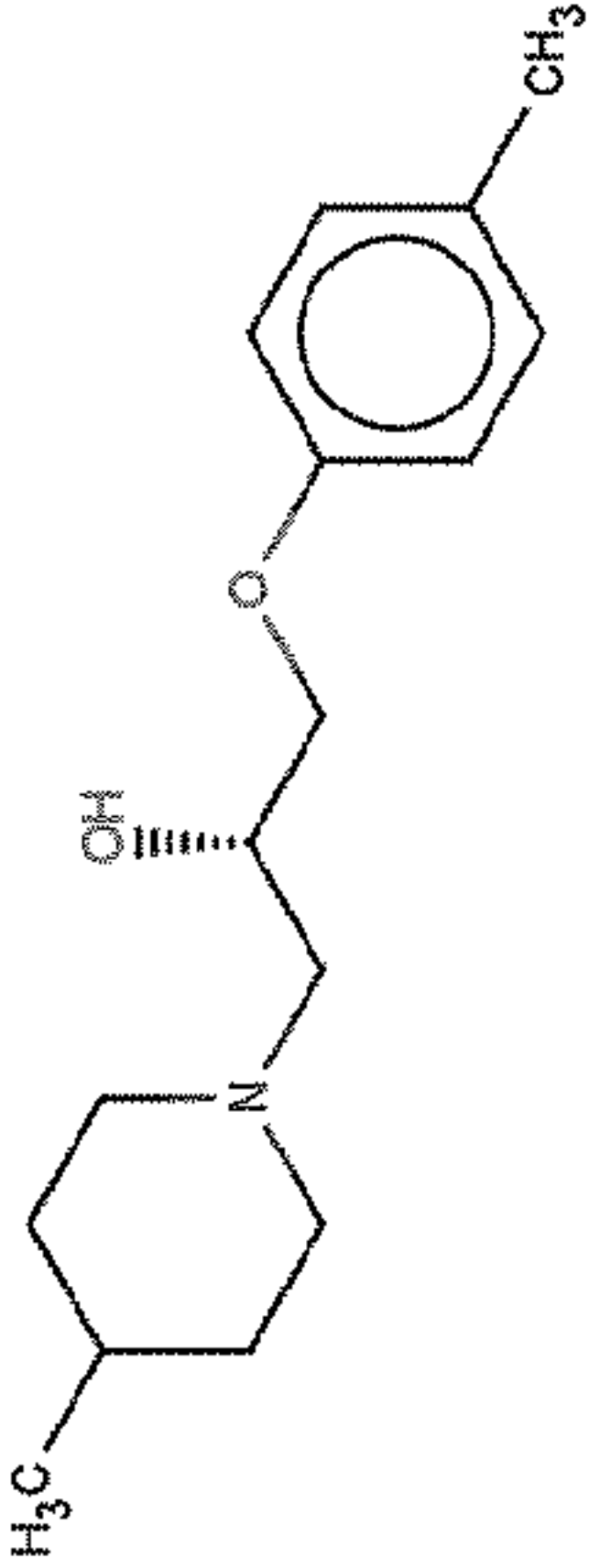
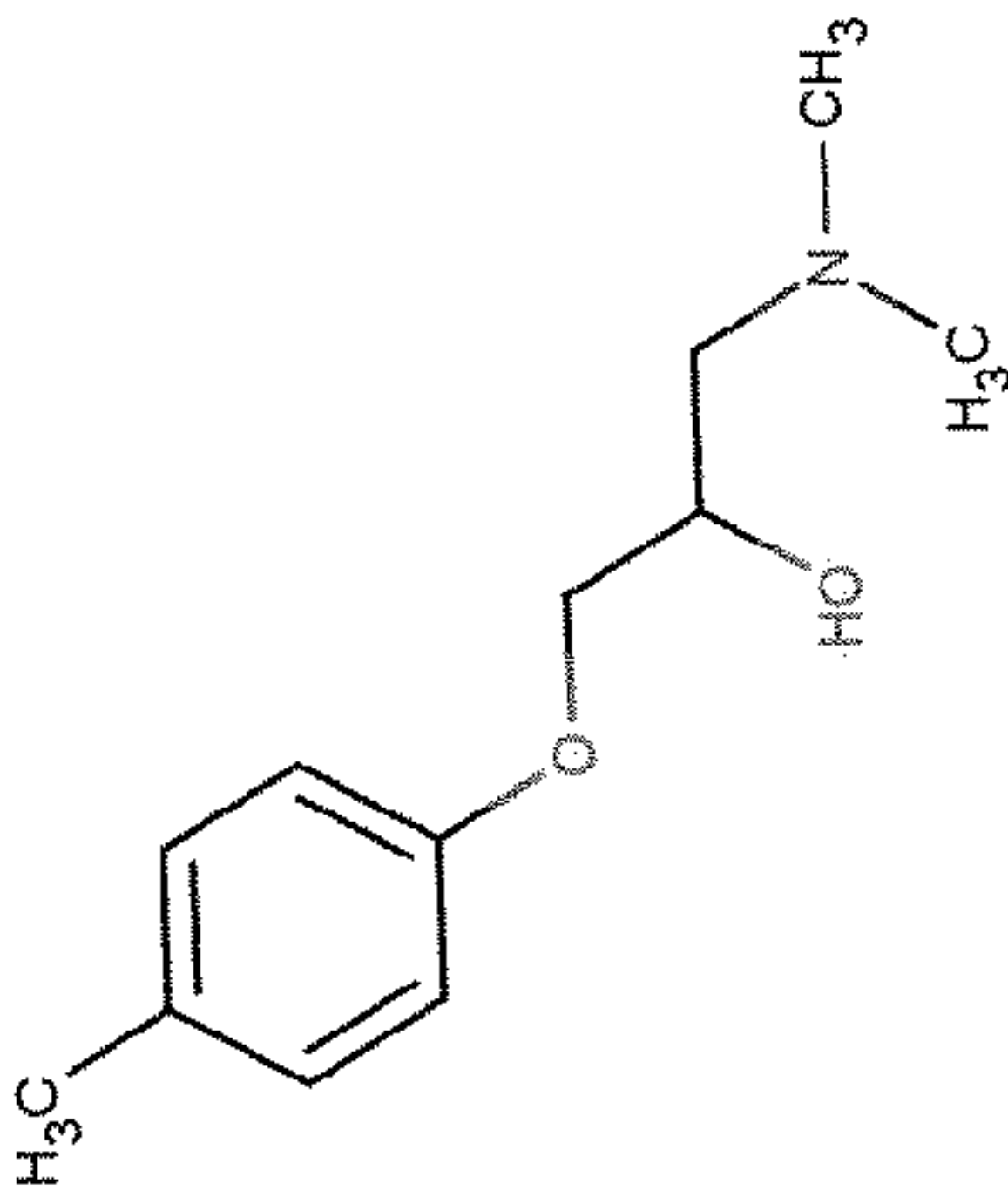
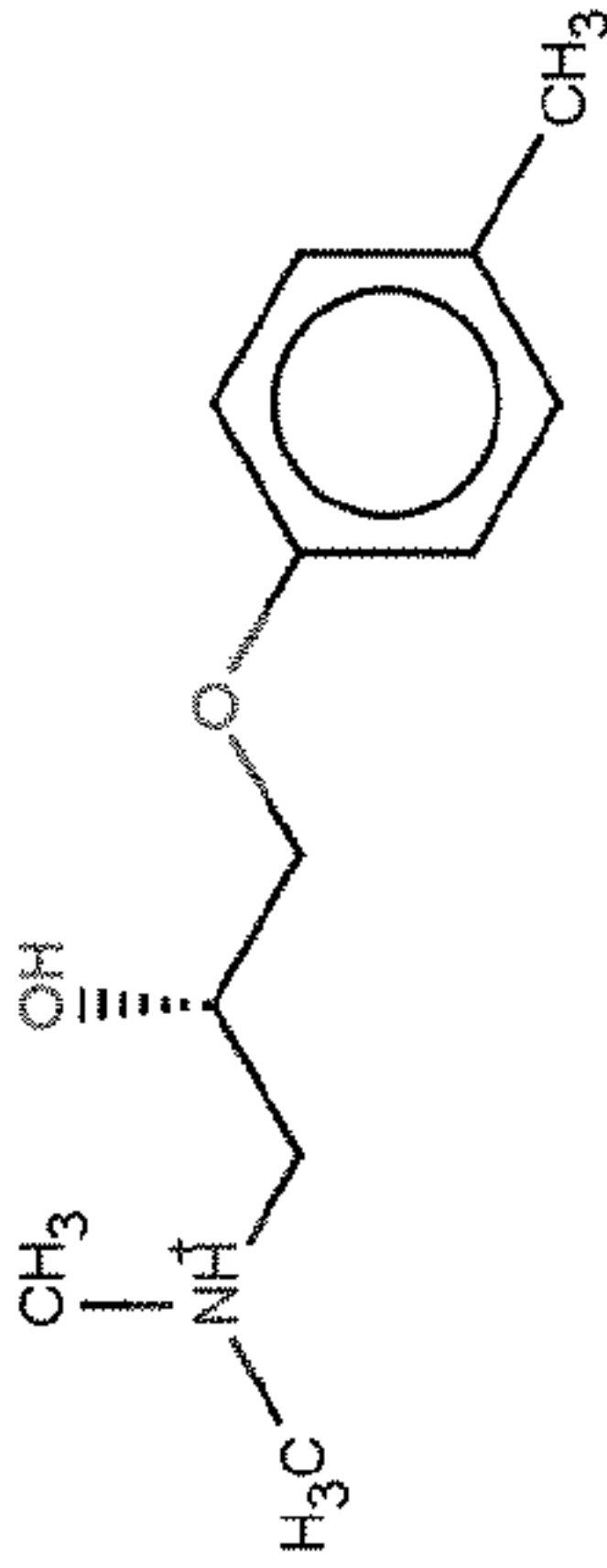
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 157		F3154-0012		F3154-0012_1 (1 of 2)
Compound 158		F3168-1703		F3168-1703_19_2 (1 of 46)
Compound 159		F3200-0073		F3200-0073_1_1 (1 of 16)

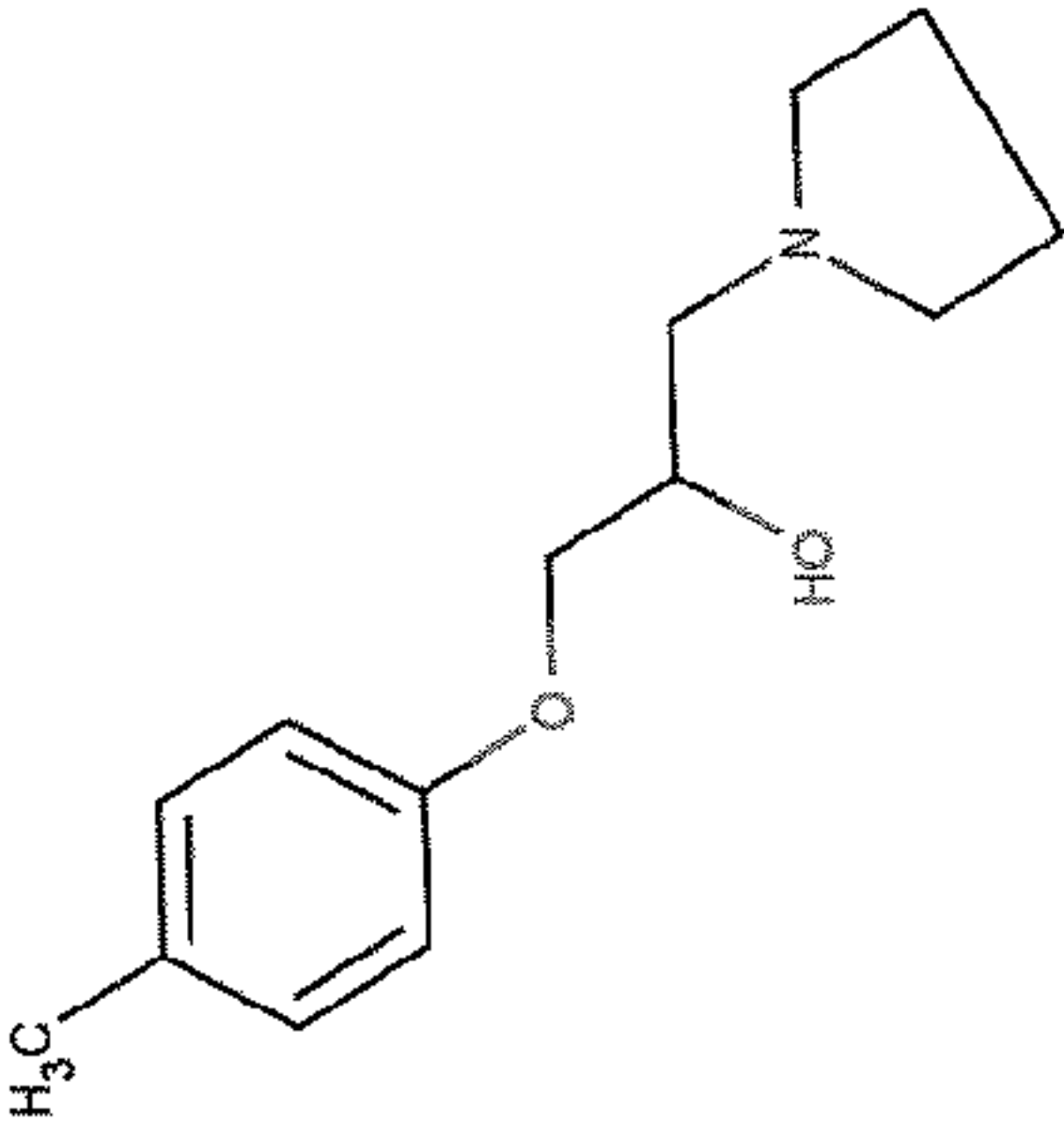
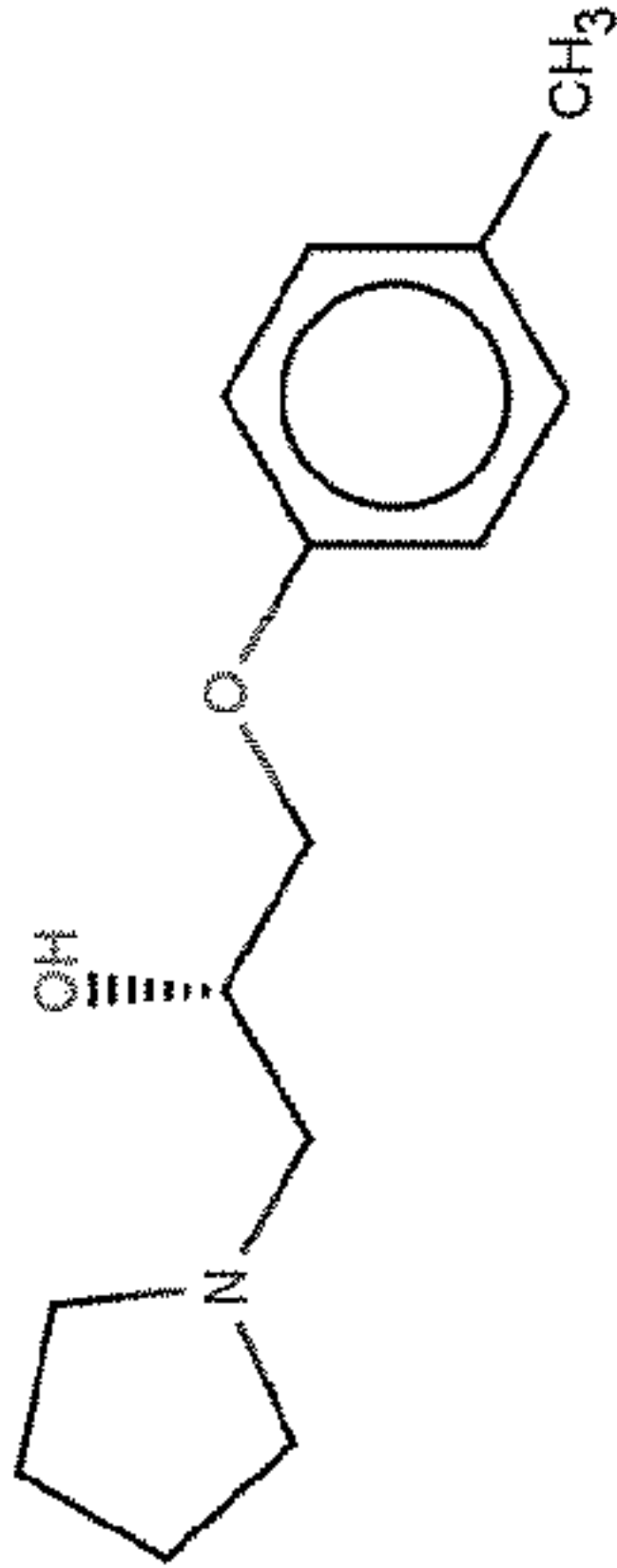
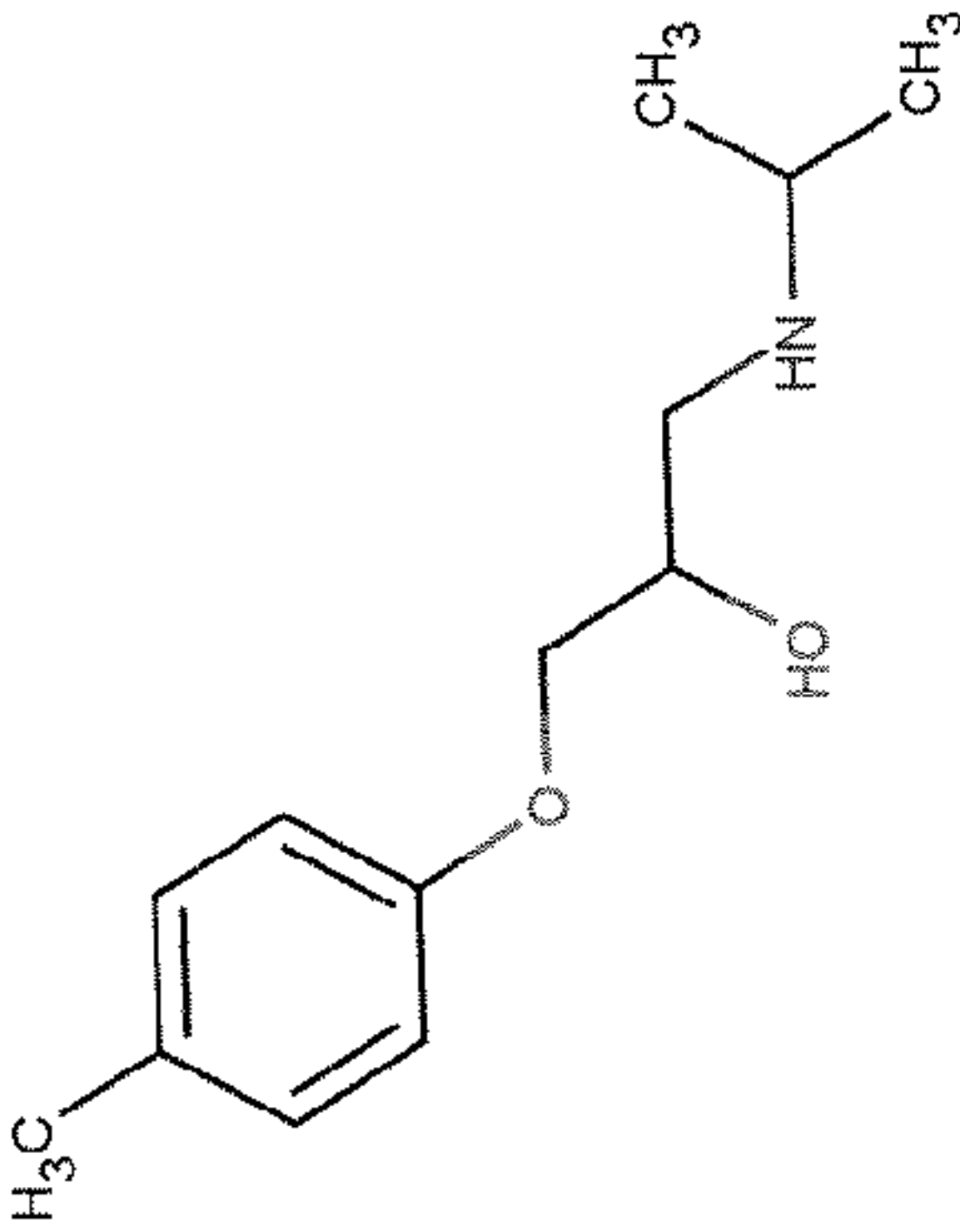
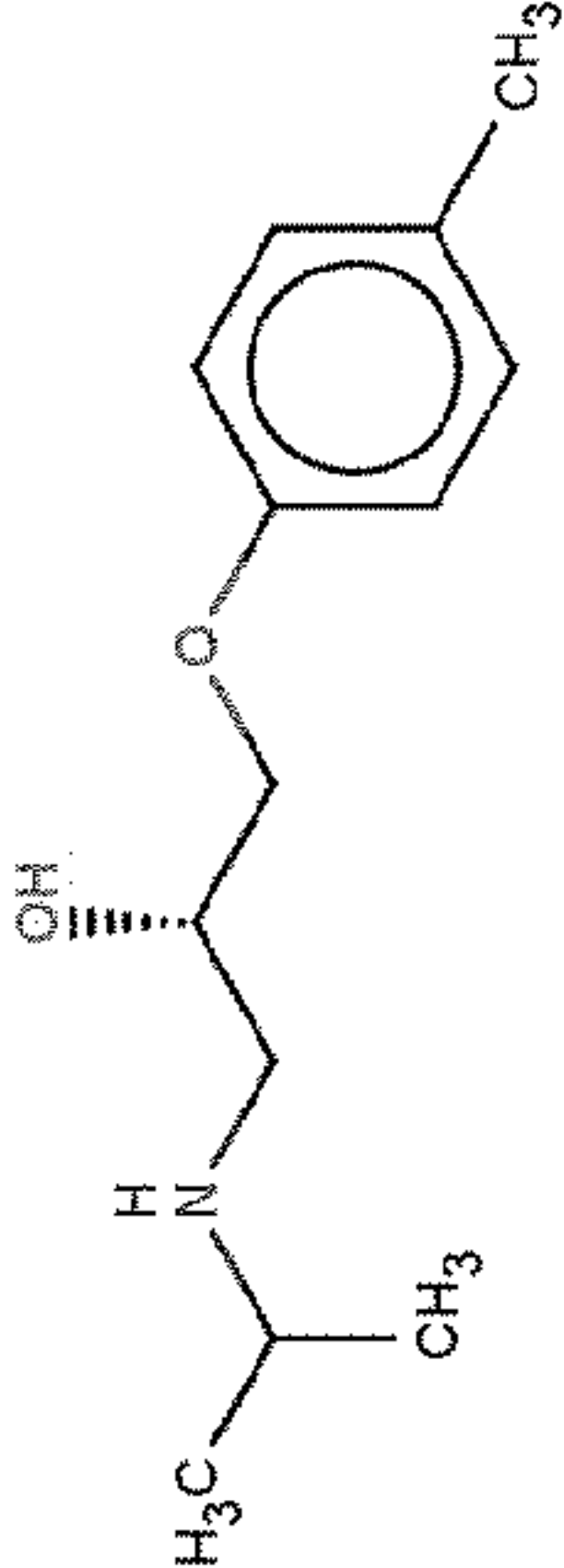
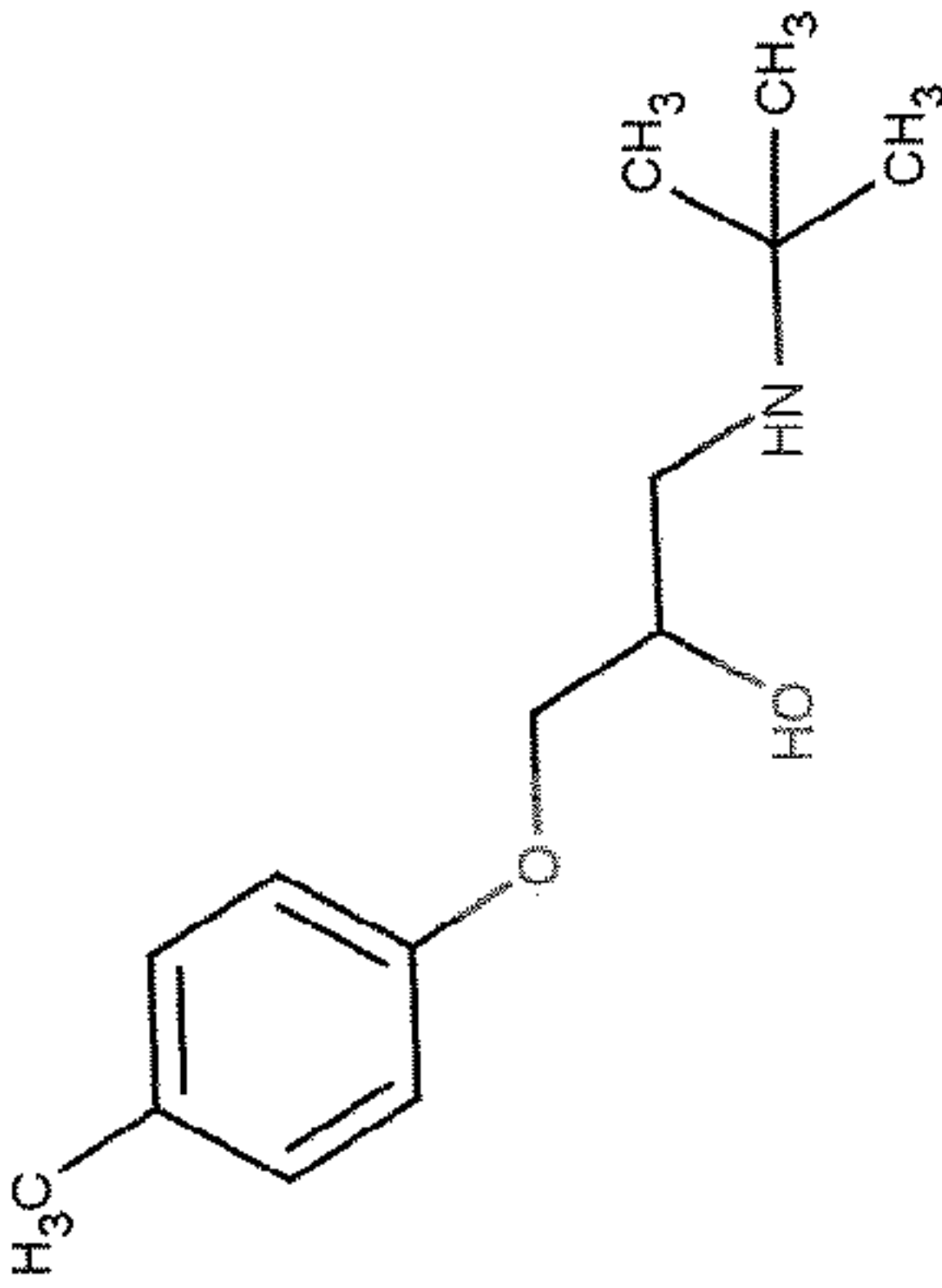
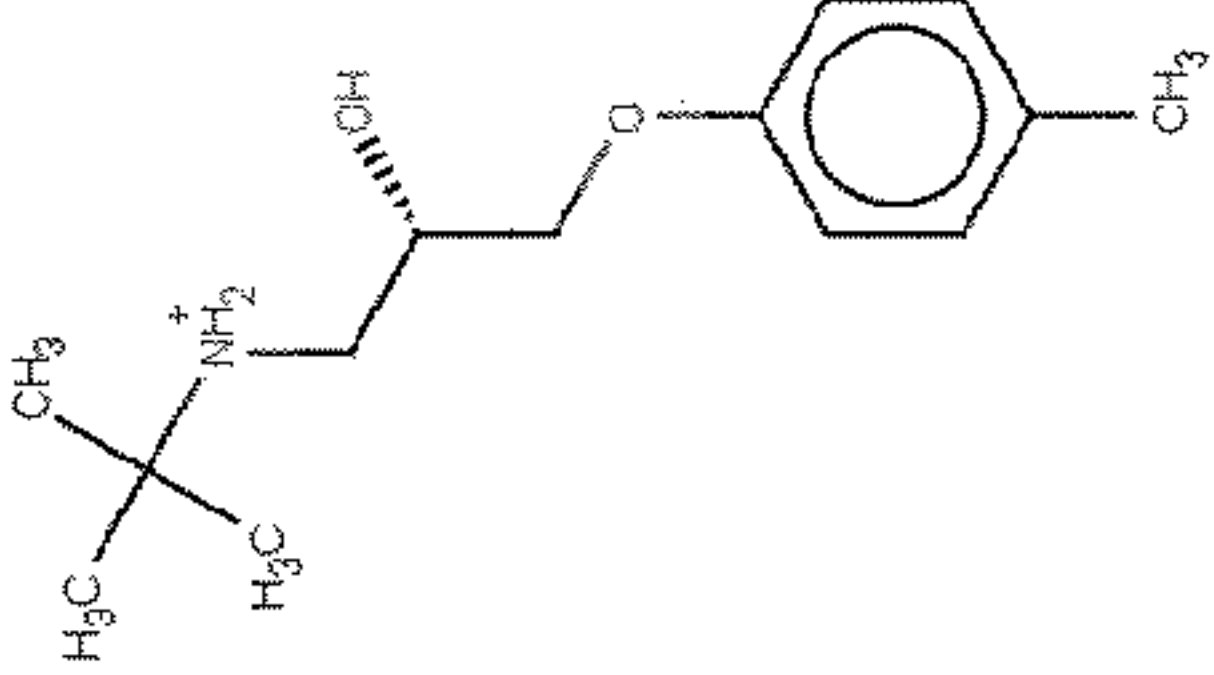
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 160		F3200-0074		F3200-0074_1_1 (1 of 16)
Compound 161		F3200-0076		F3200-0076_1_17 (1 of 53)
Compound 162		F3219-0003		F3219-0003_23_1 (1 of 27)

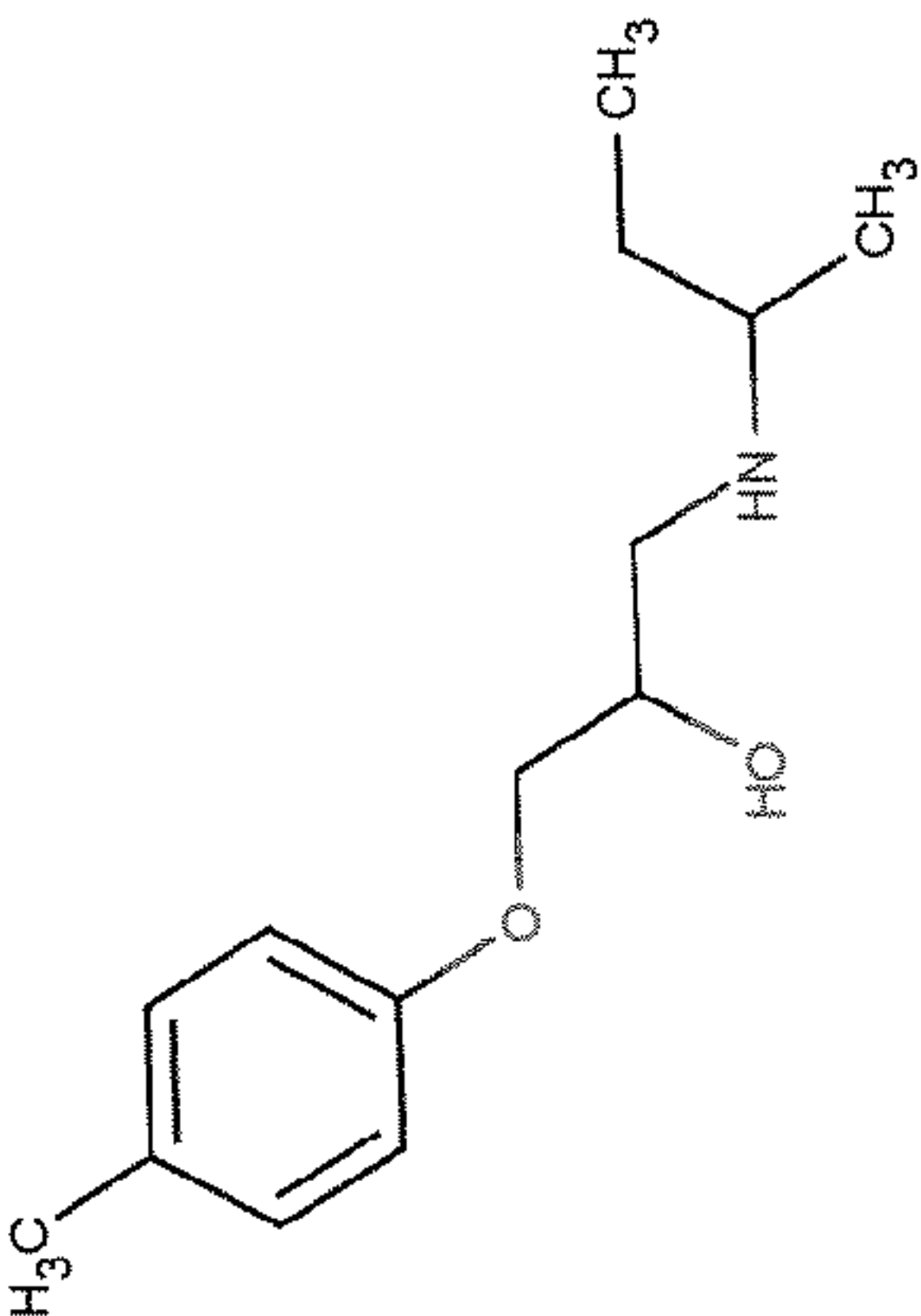
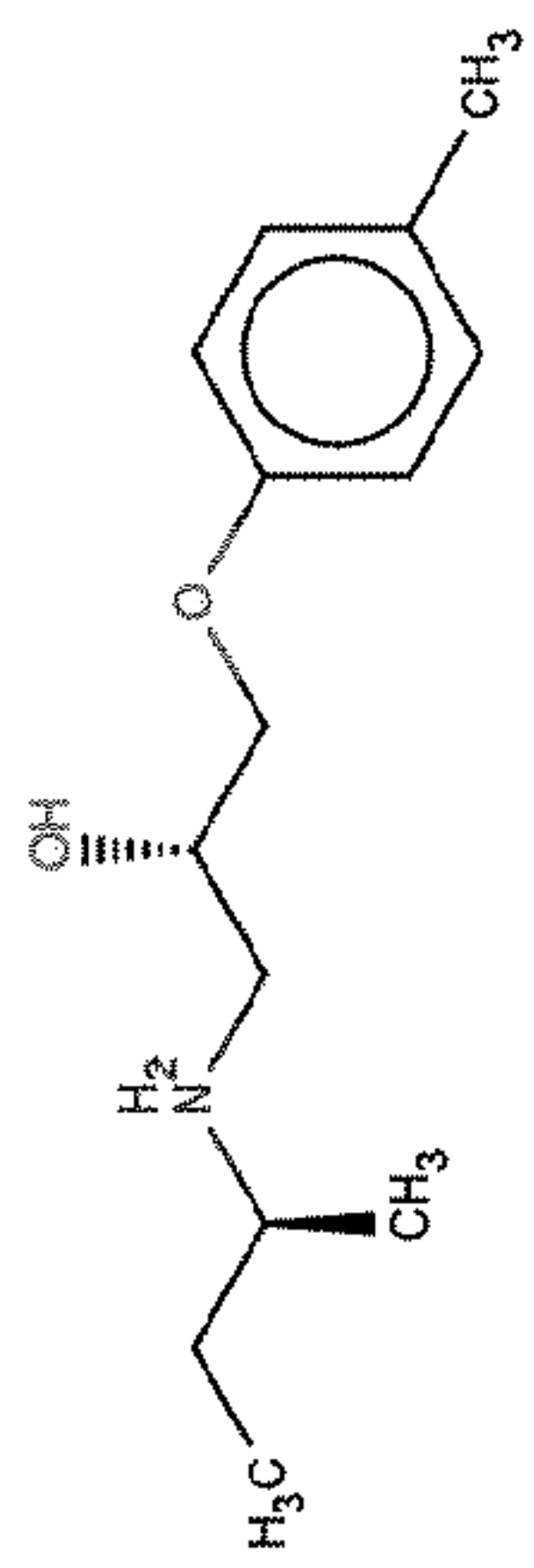
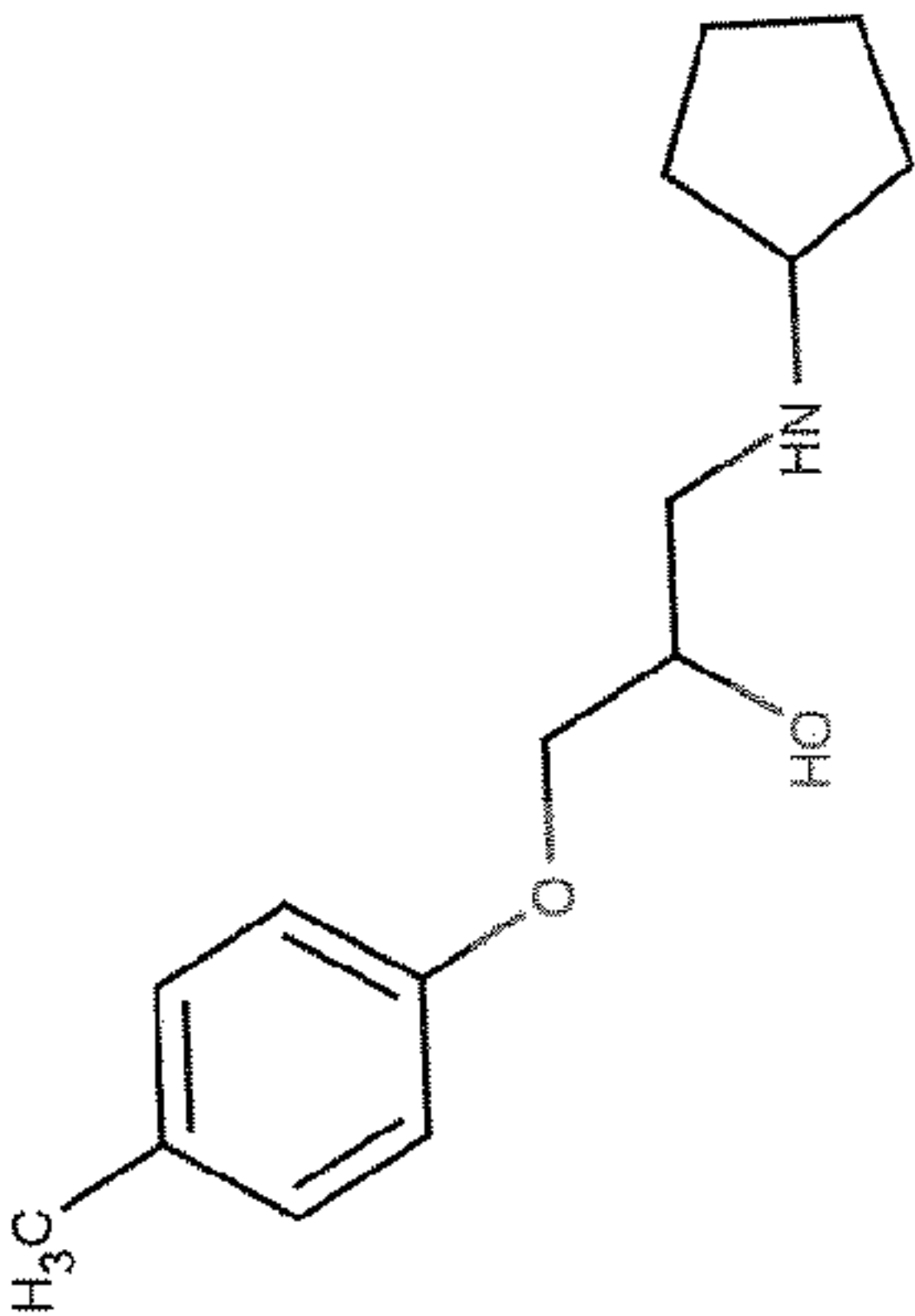
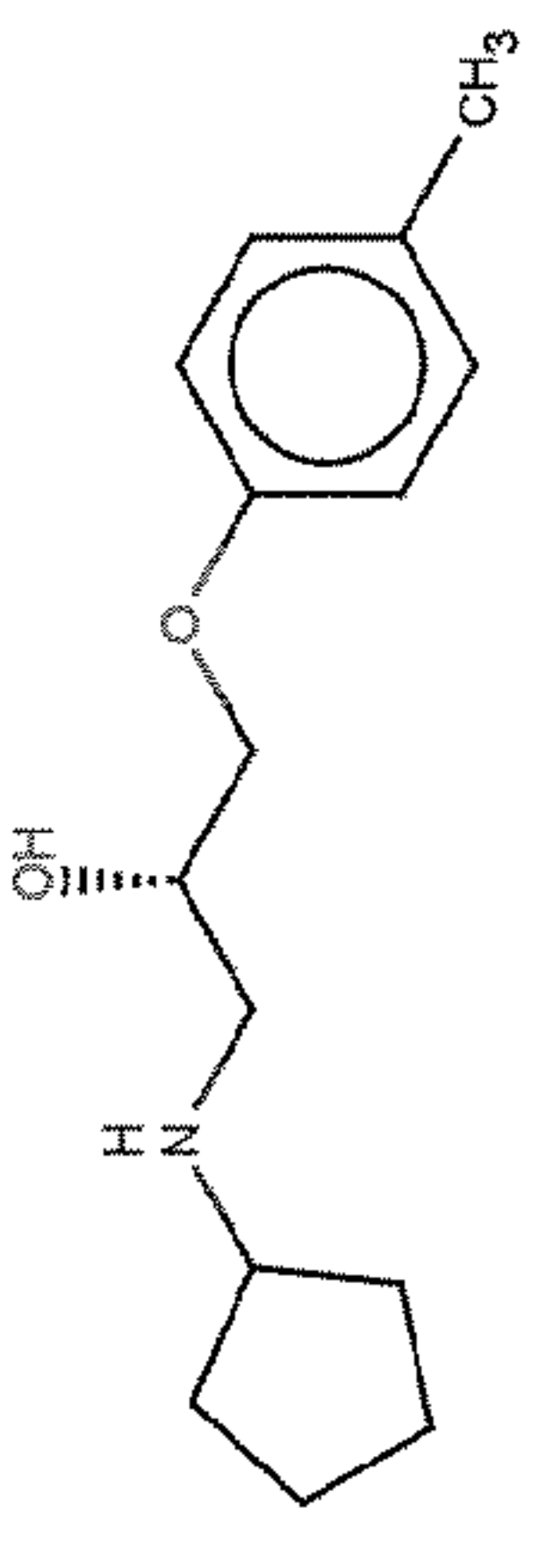
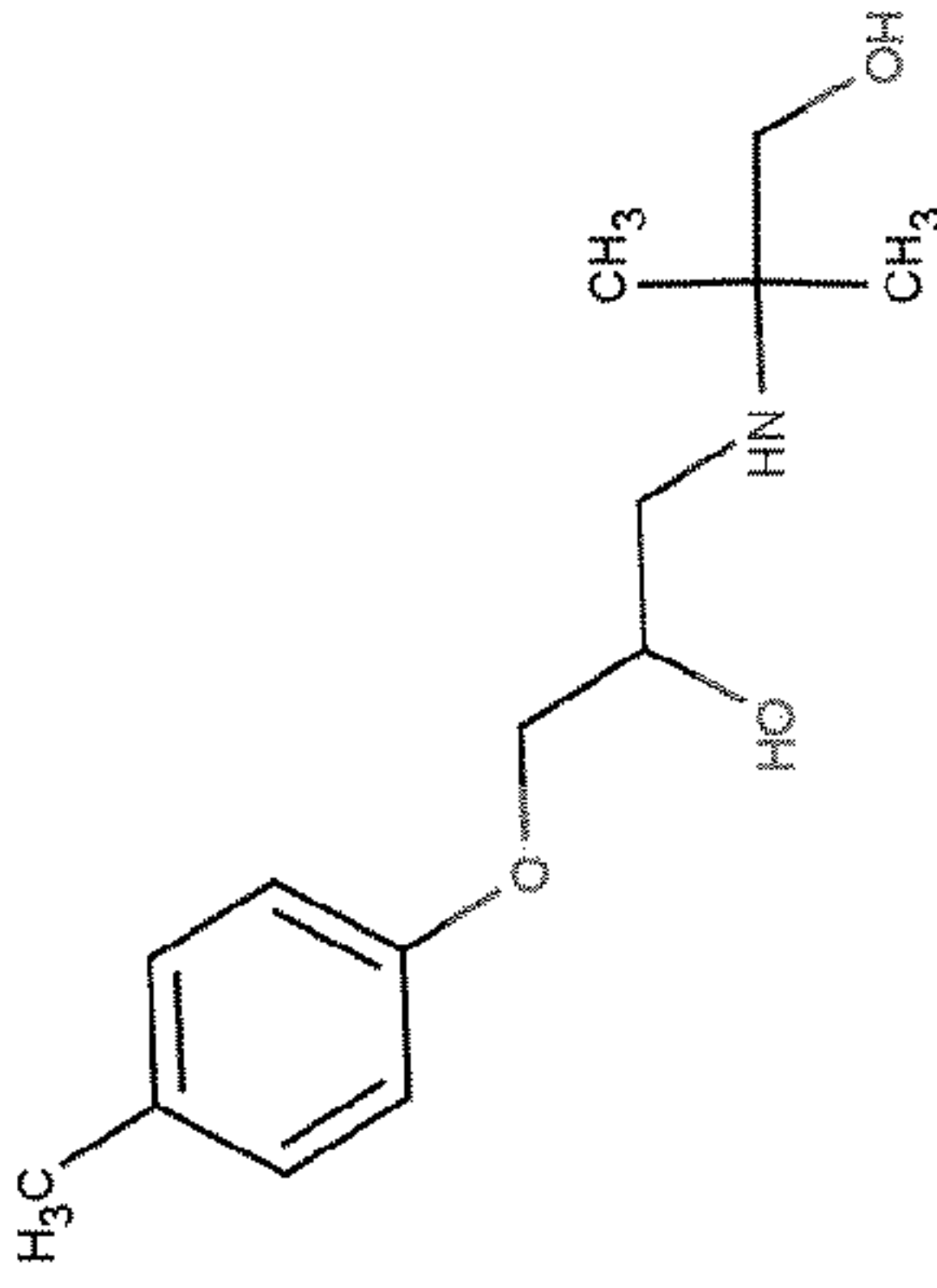
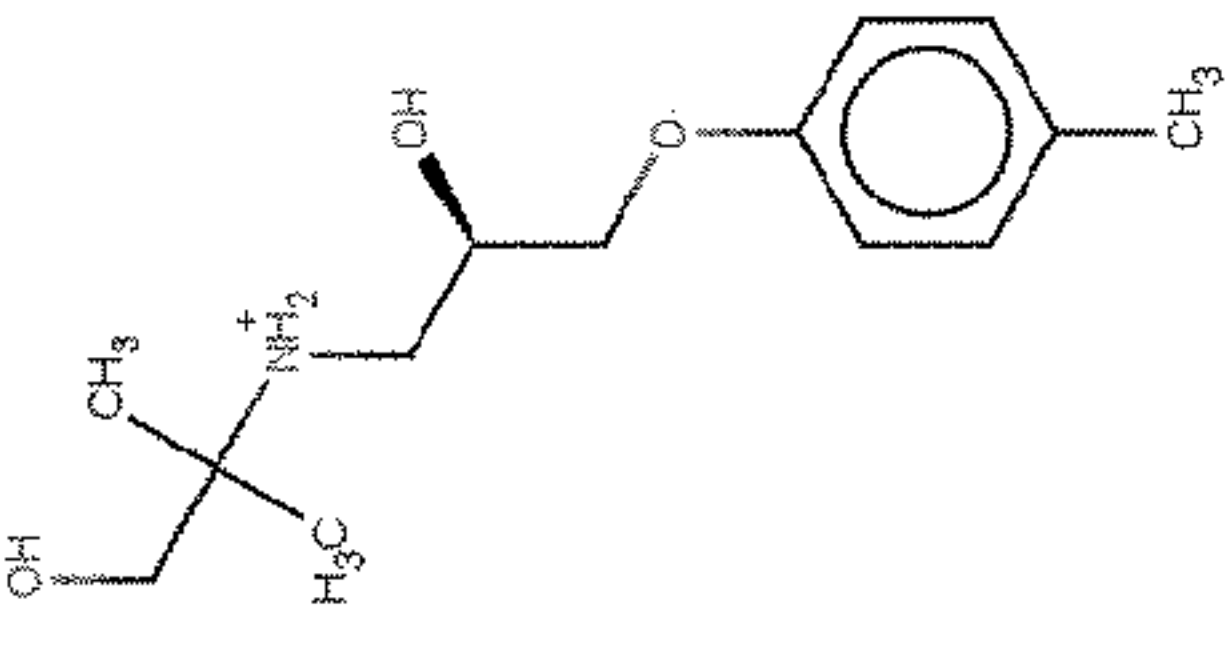
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 163		F3225-8622		F3225-8622_3_1 (1 of 4)
Compound 164		F3230-0054		F3230-0054_1 (1 of 1)
Compound 165		F3250-0673		F3250-0673_1 (1 of 4)

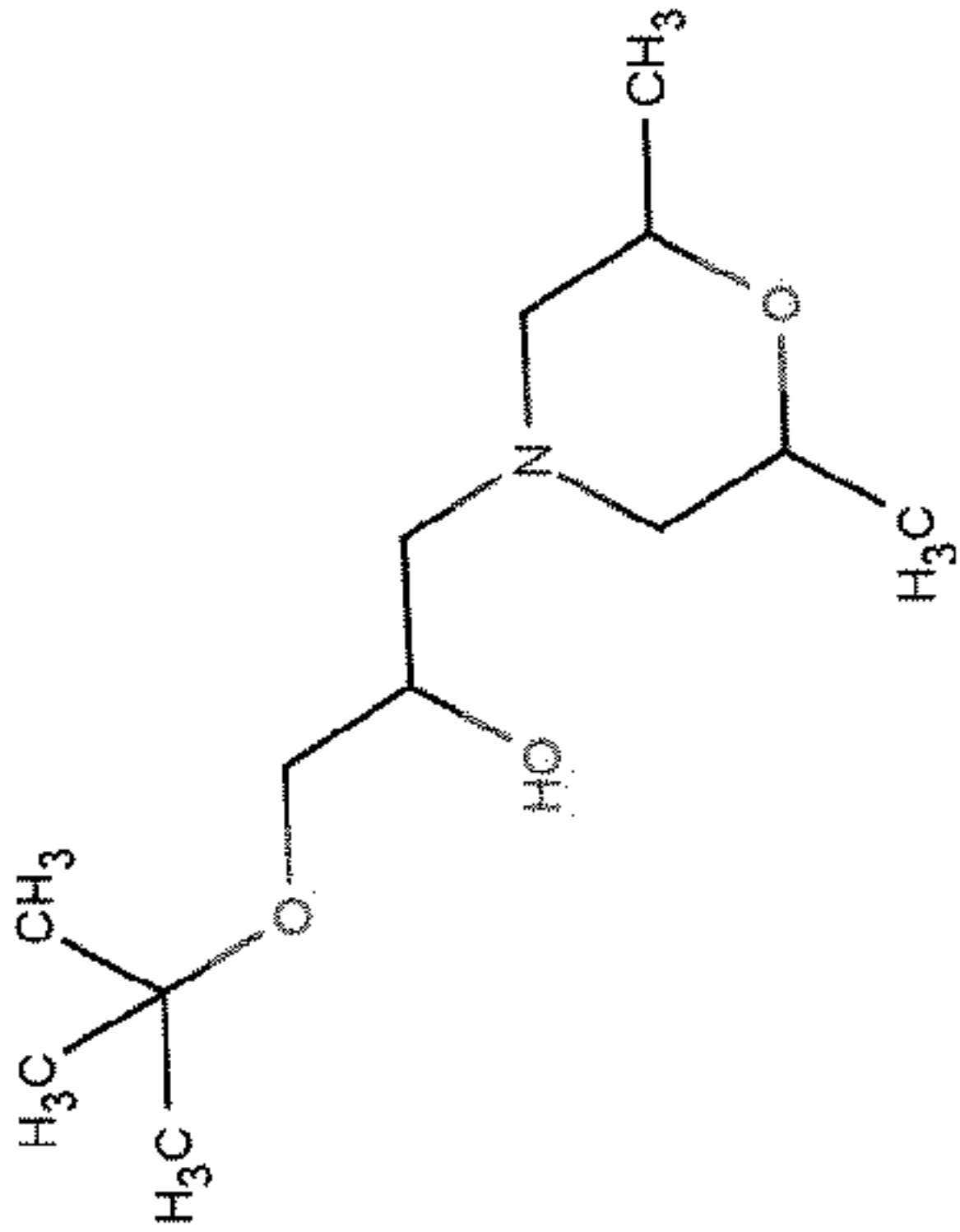
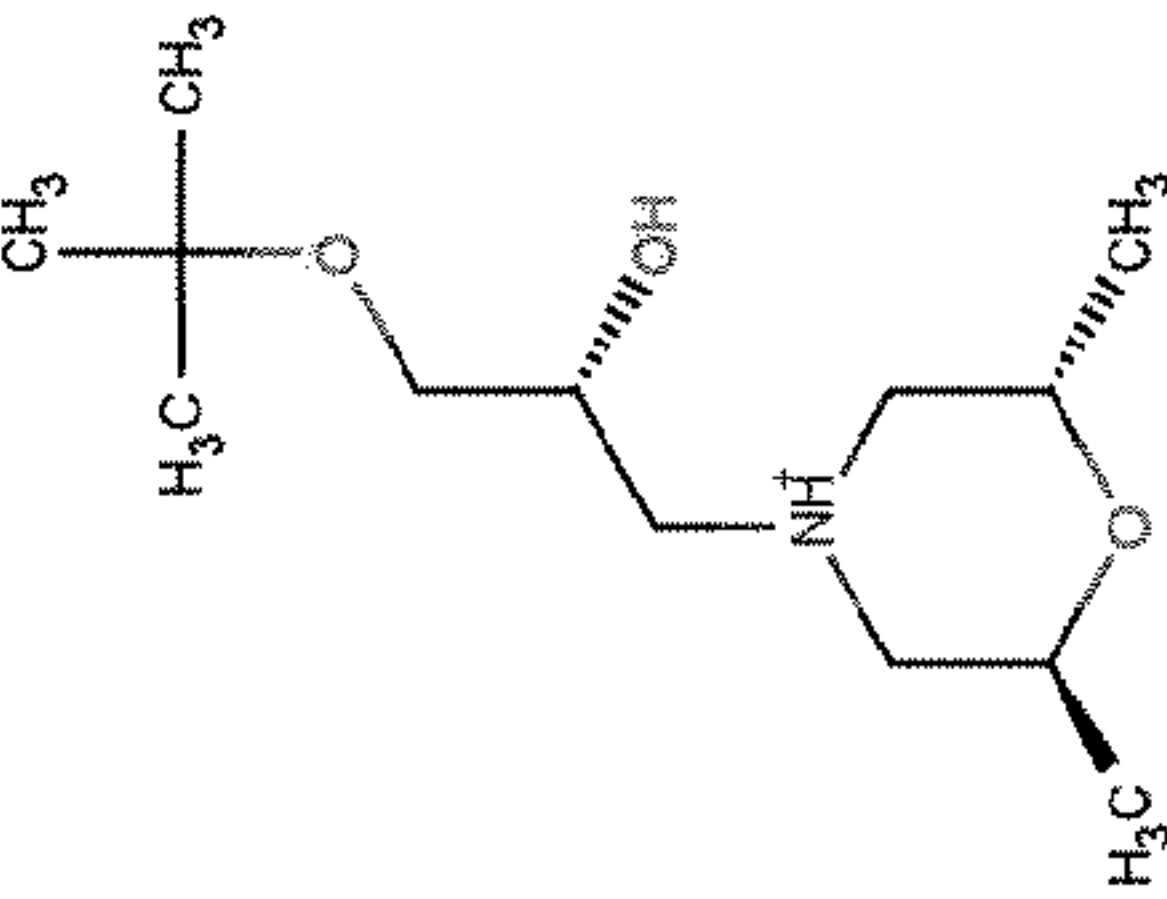
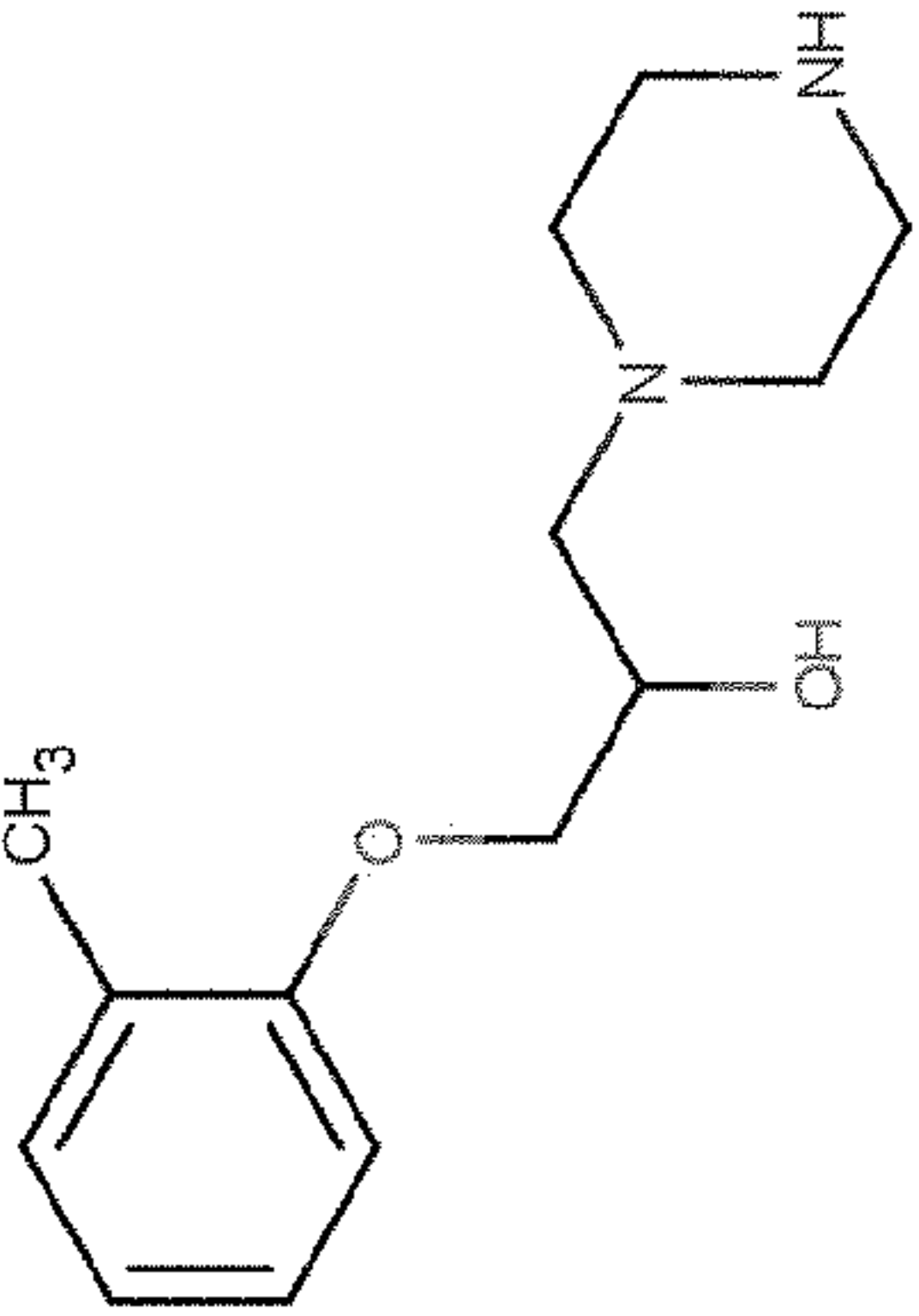
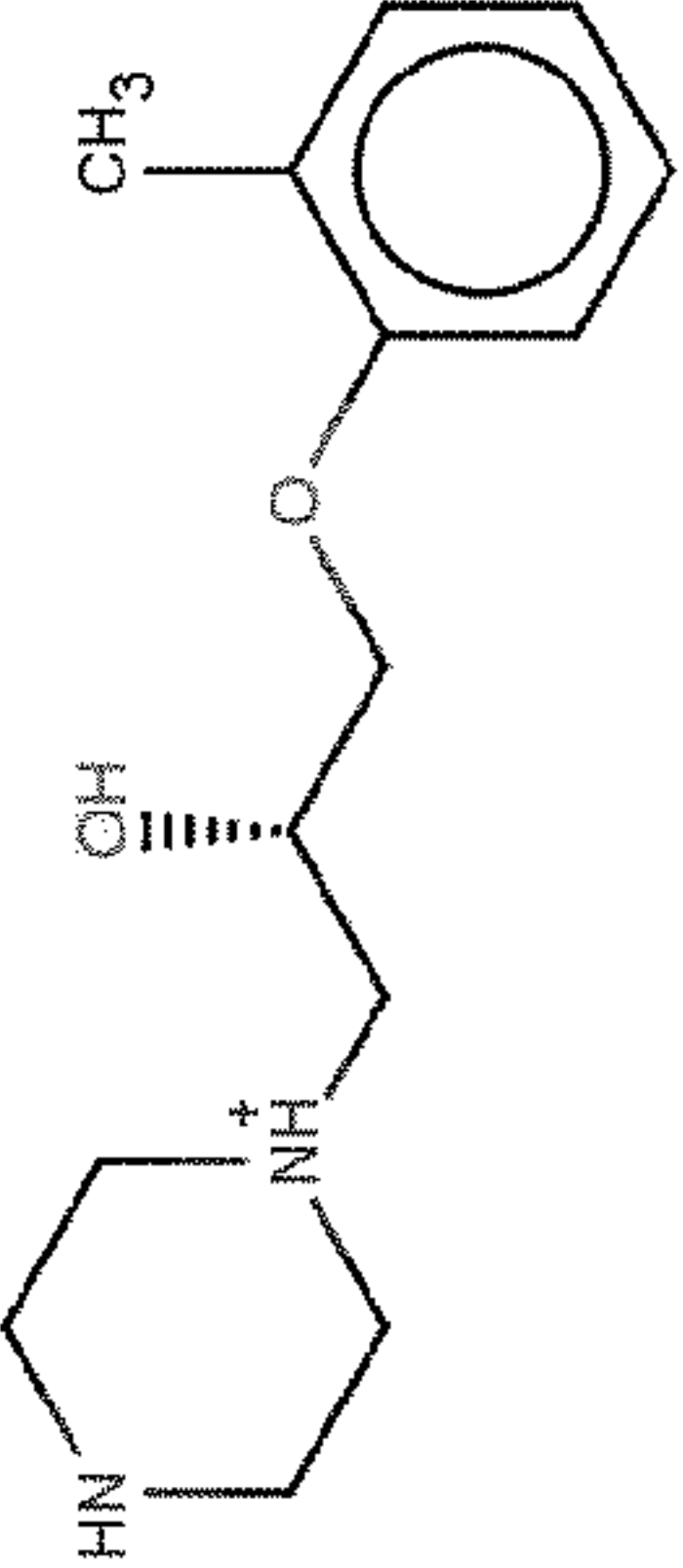
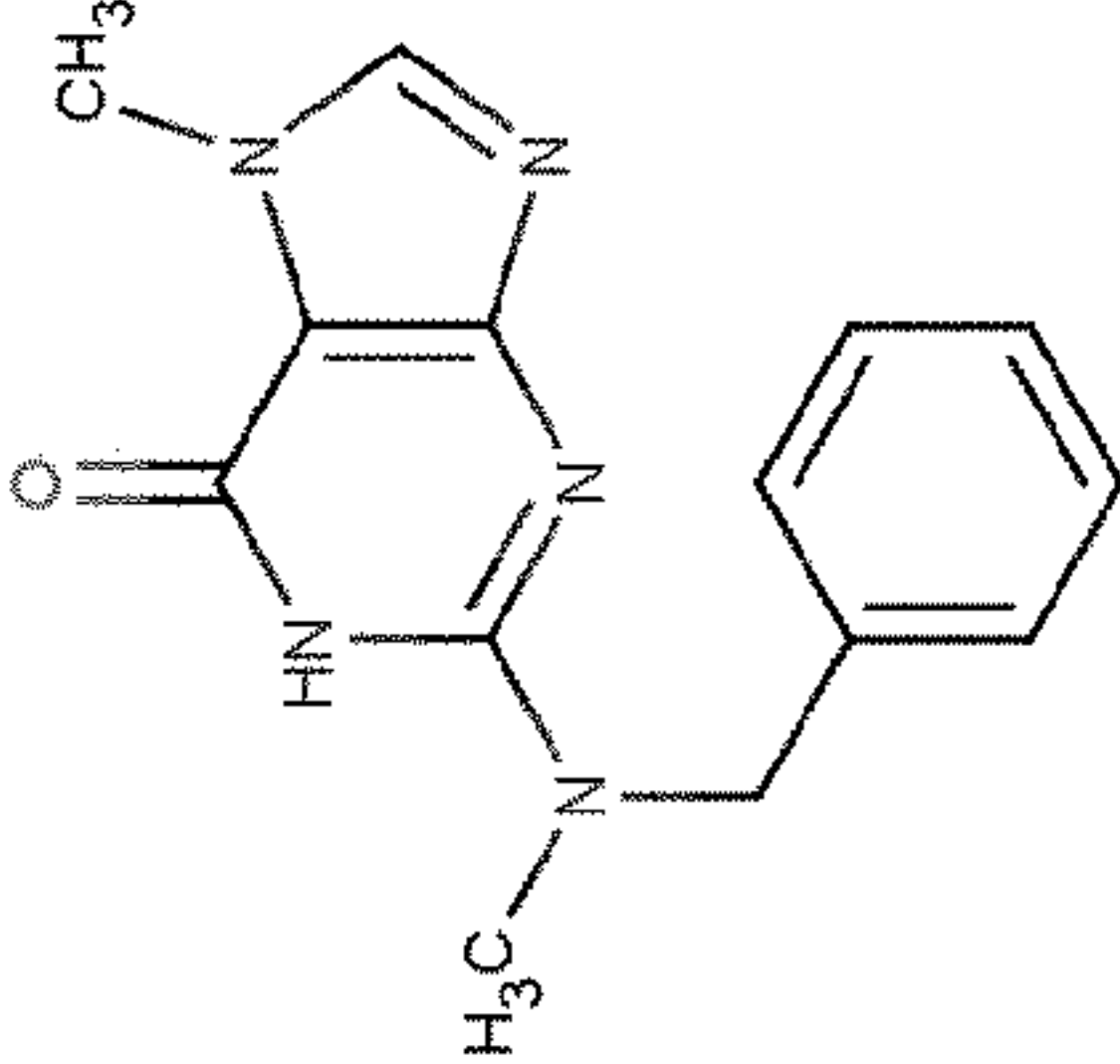
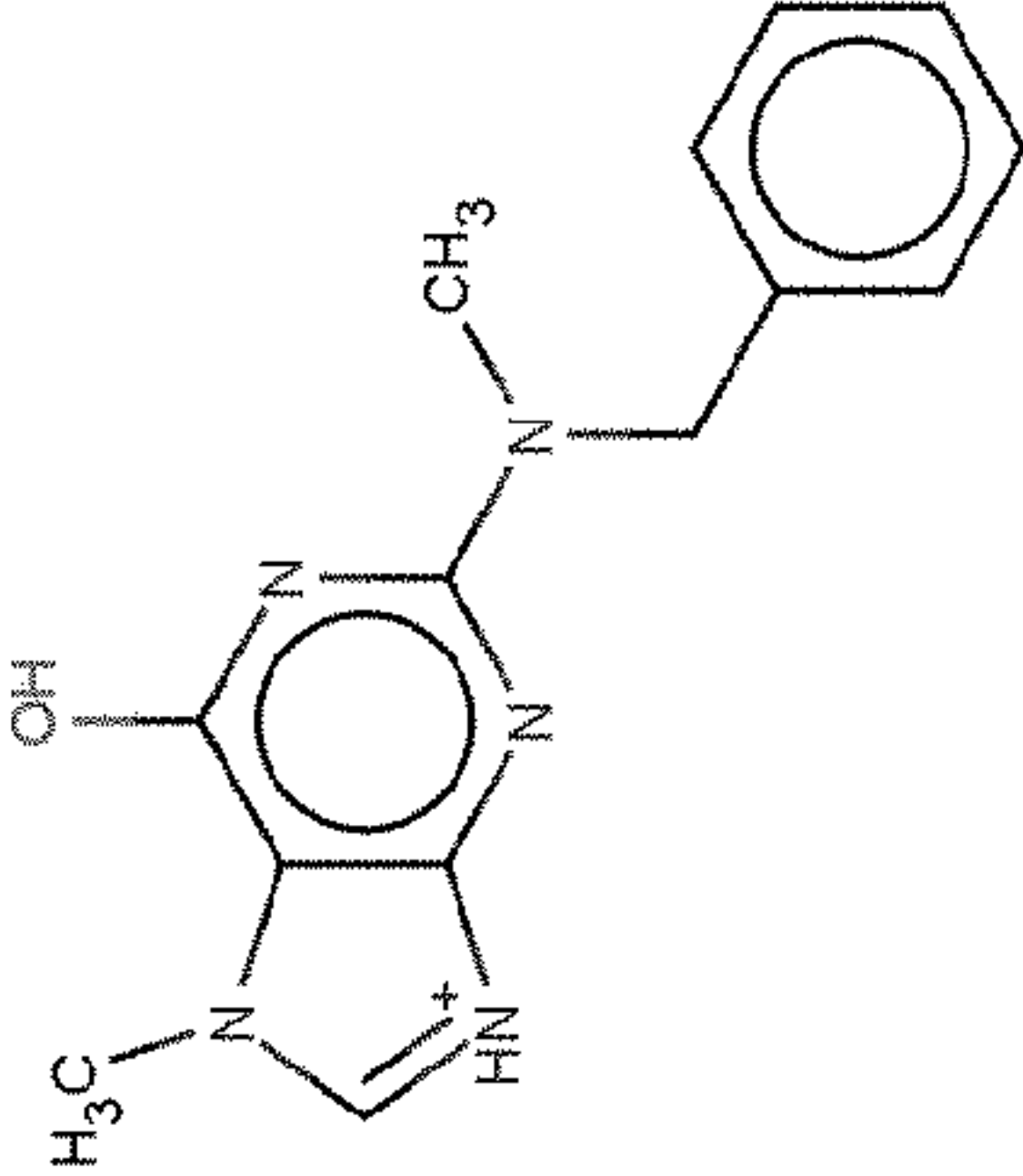
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 166		F3250-0686		F3250-0686_2_2 (1 of 14)
Compound 167		F3250-0689		F3250-0689_1 (1 of 4)
Compound 168		F3250-0696		F3250-0696_1 (1 of 4)

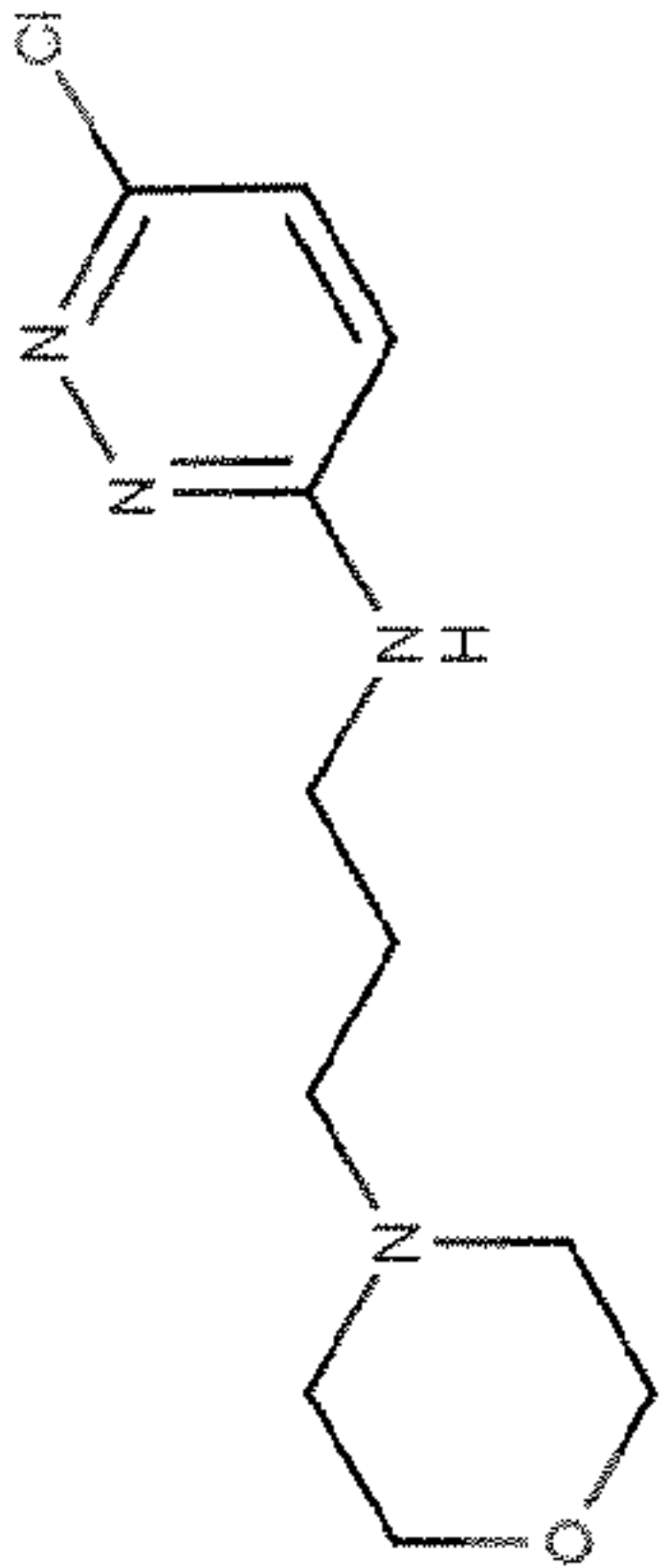
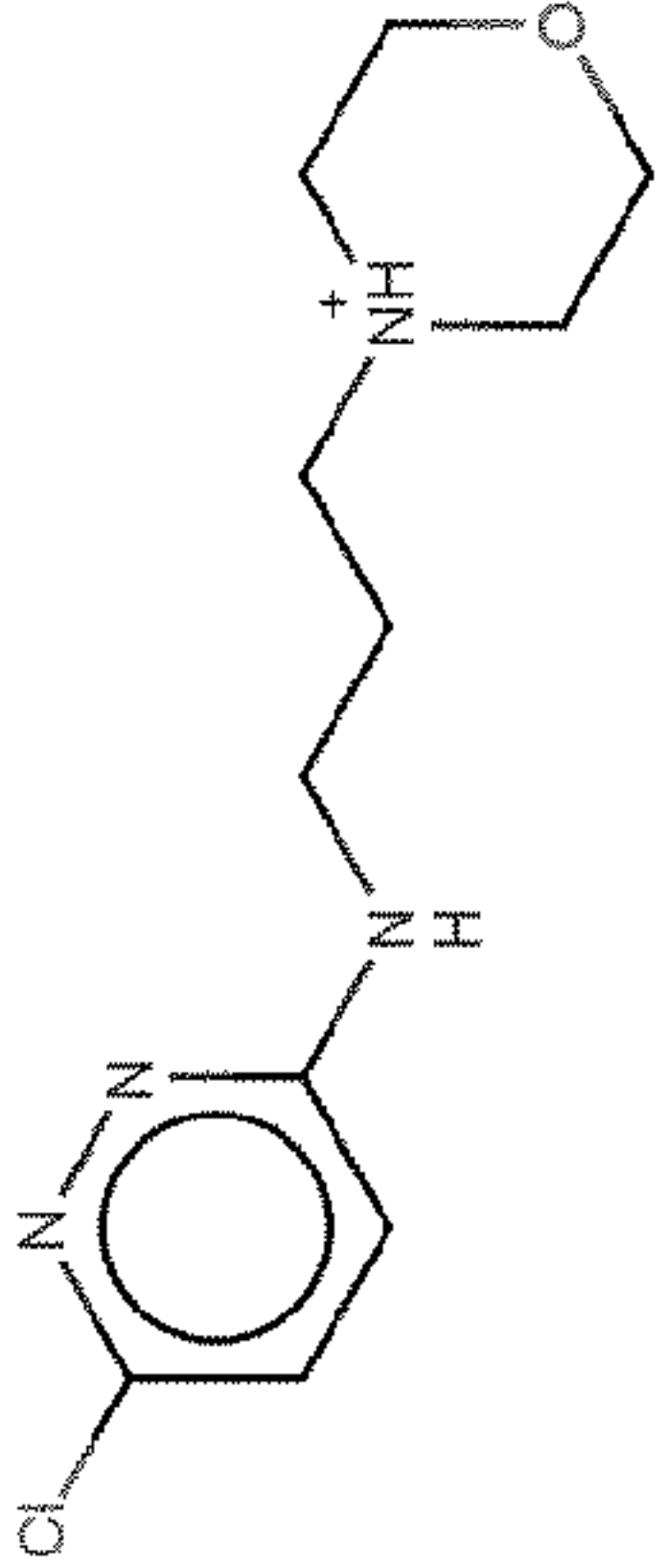
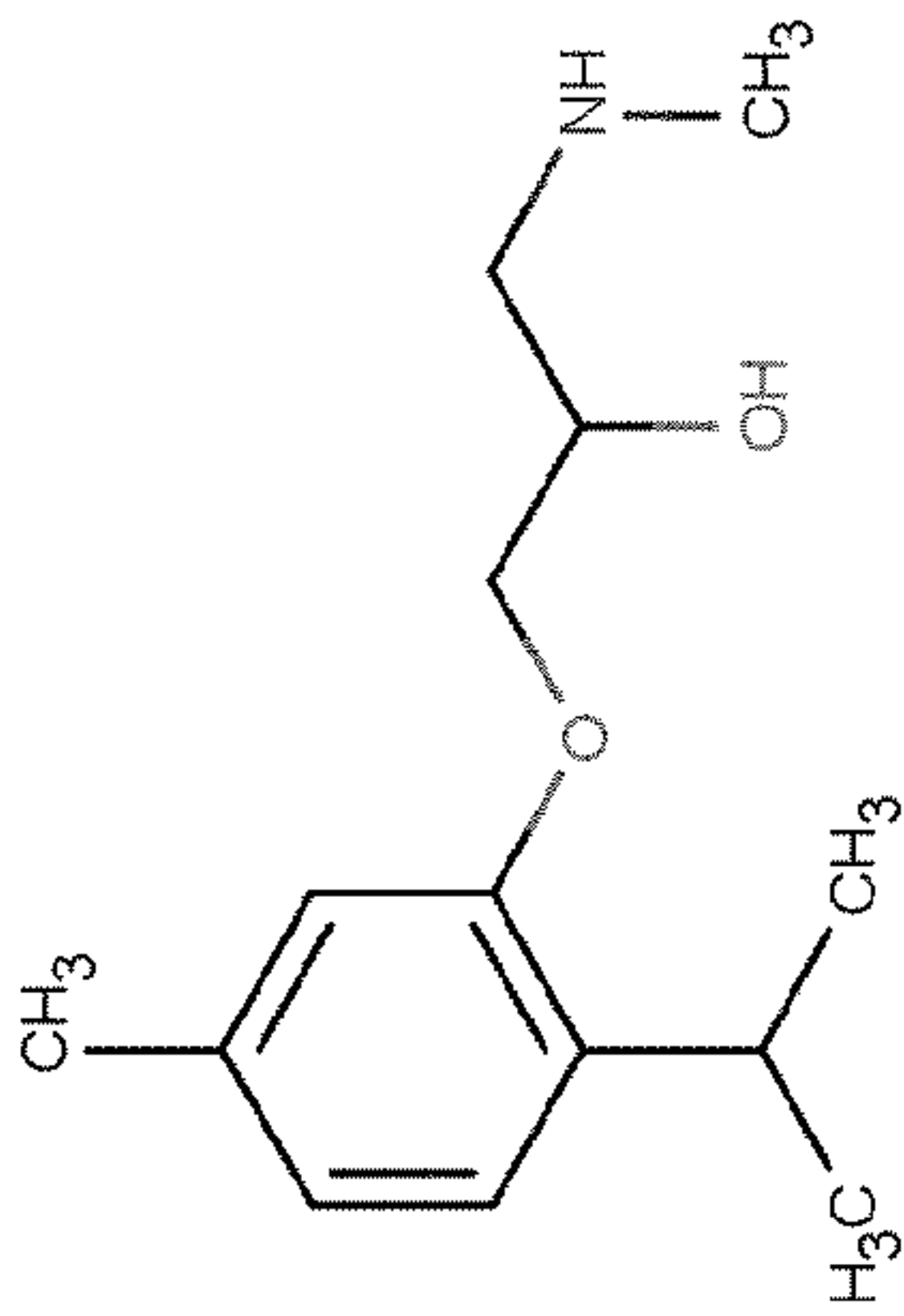
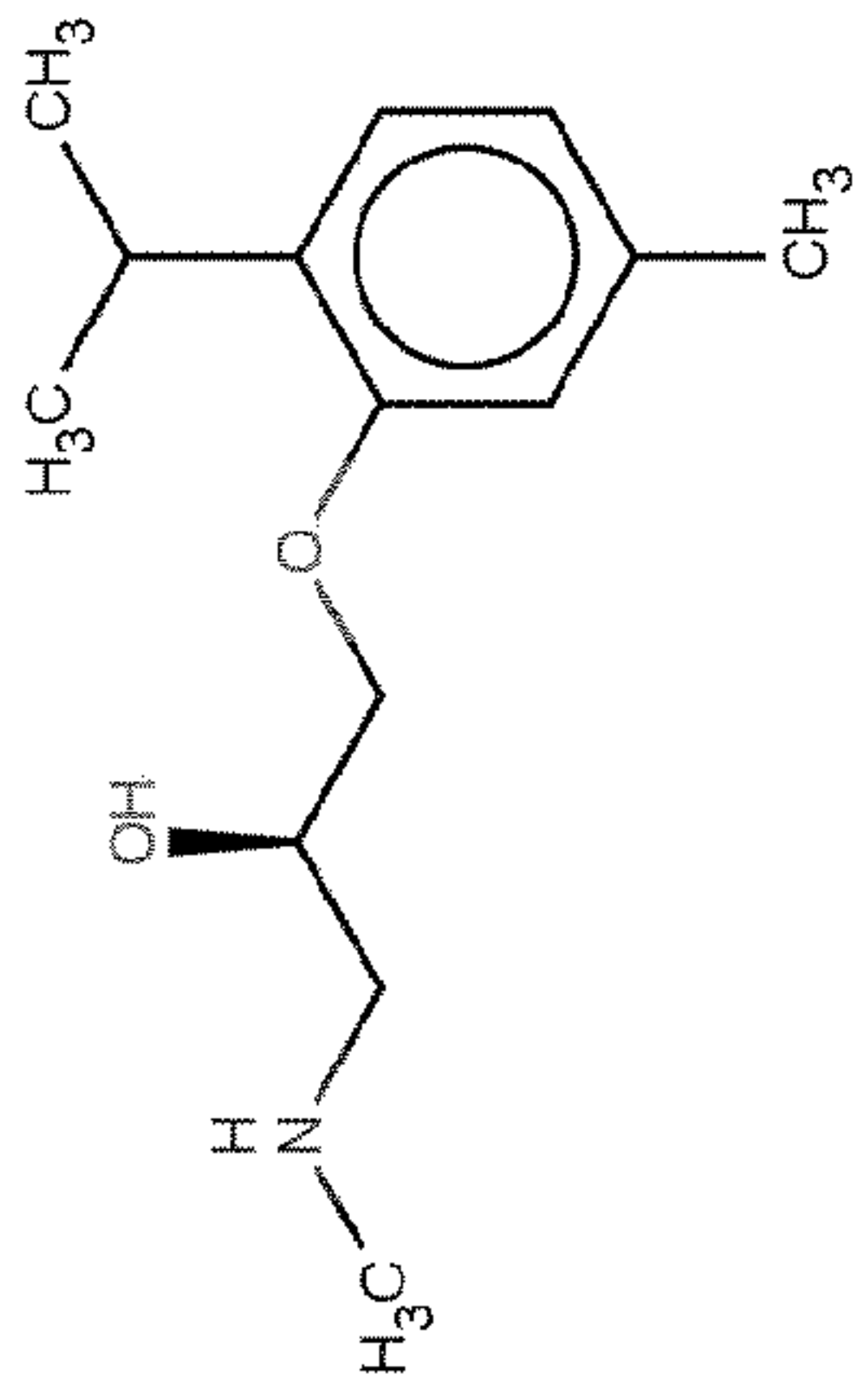
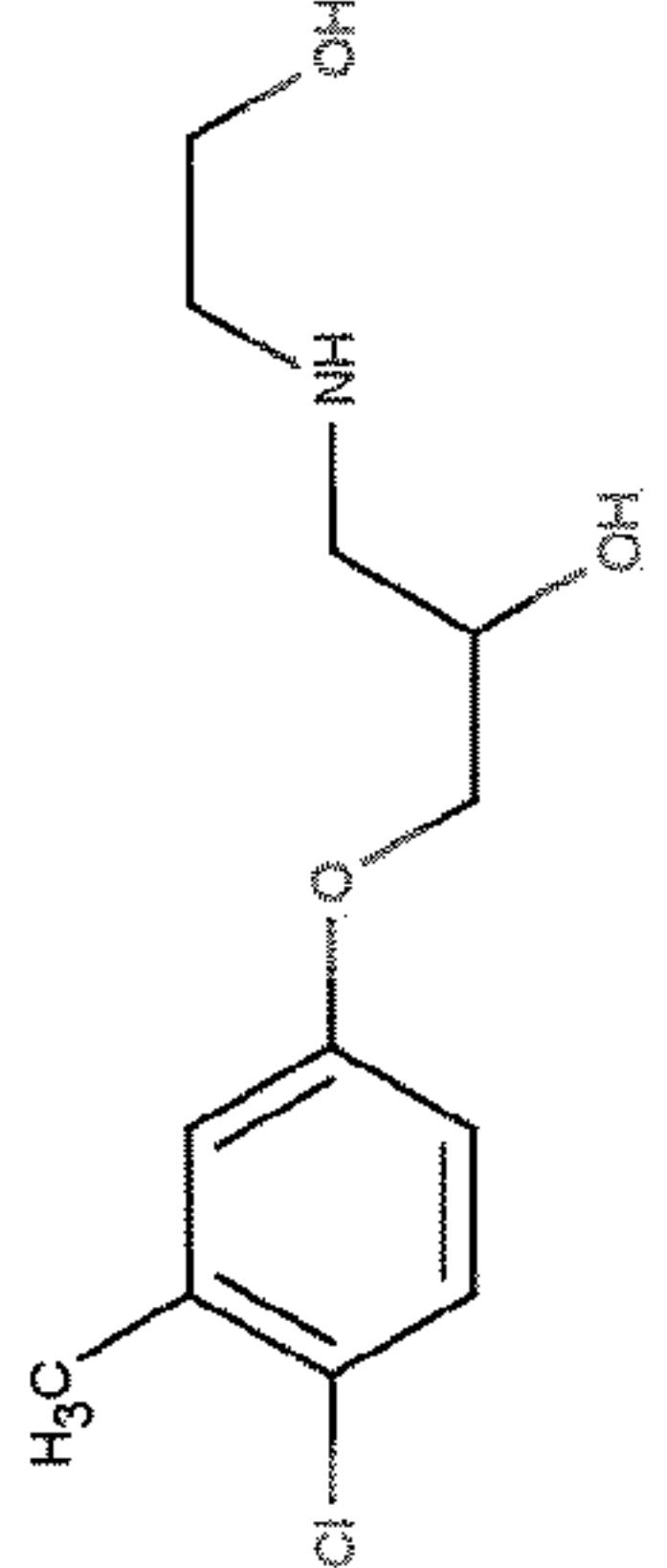
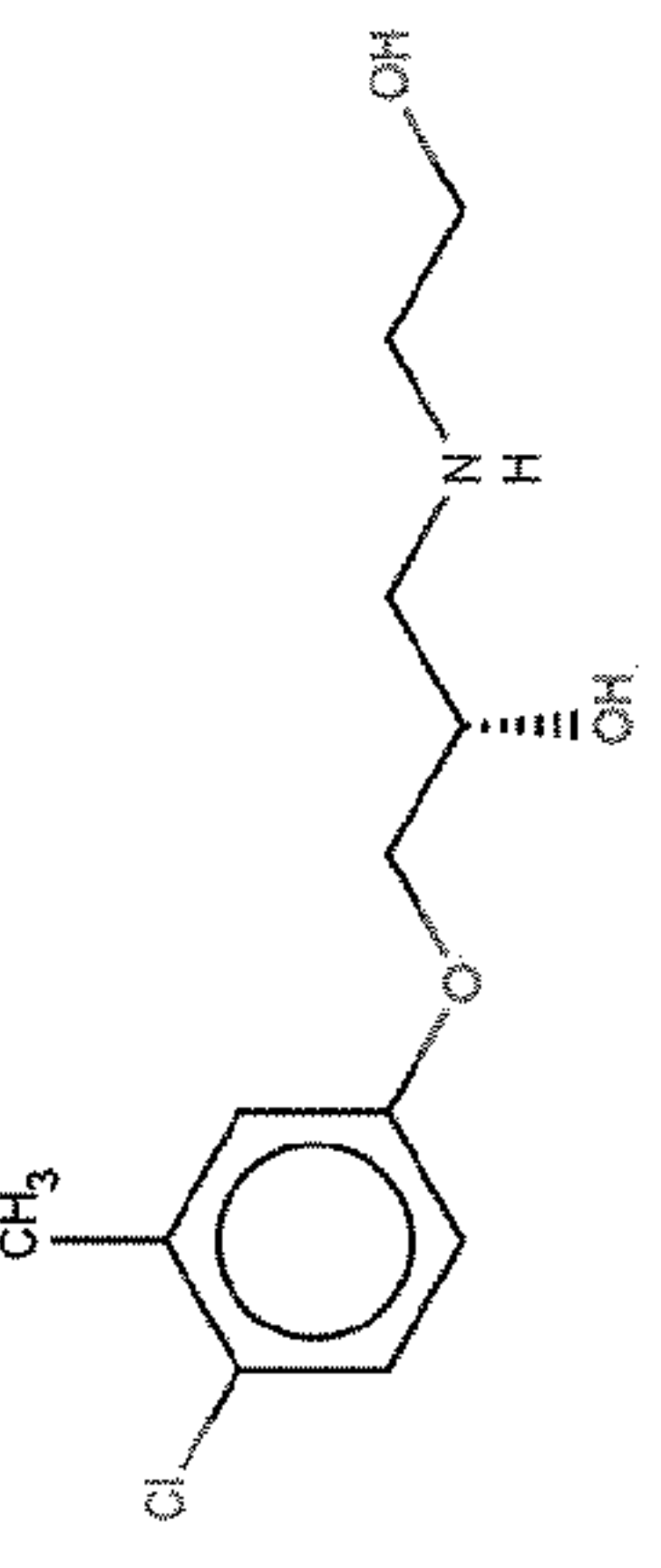
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 169		F3250-0733		F3250-0733_1_1 (1 of 4)
Compound 170		F3254-0053		F3254-0053_1_1 (1 of 4)
Compound 171		F3254-0054		F3254-0054_3 (1 of 8)

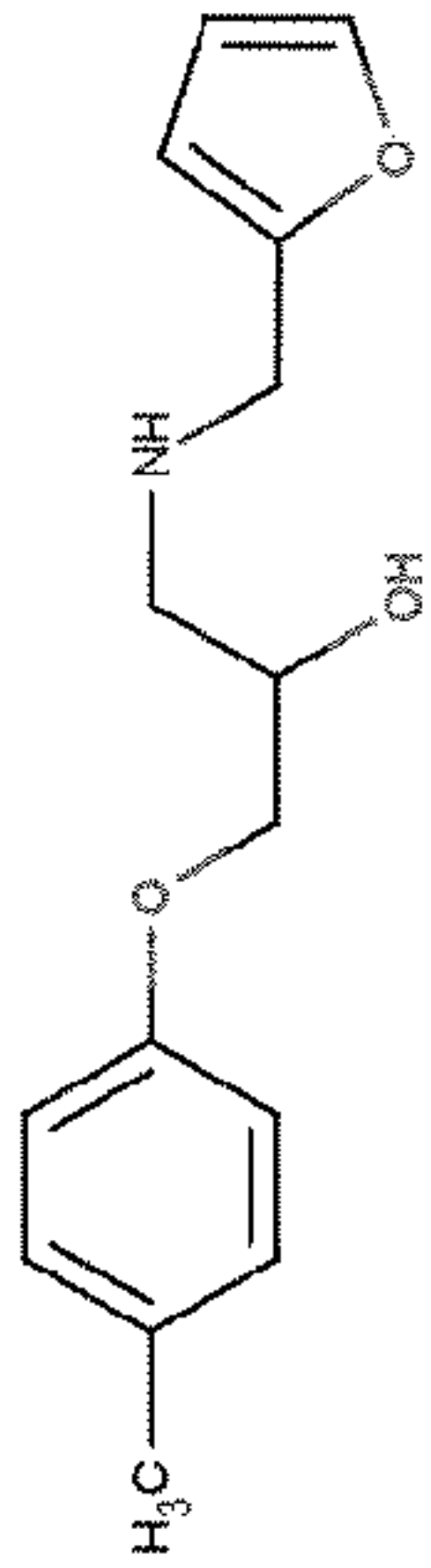
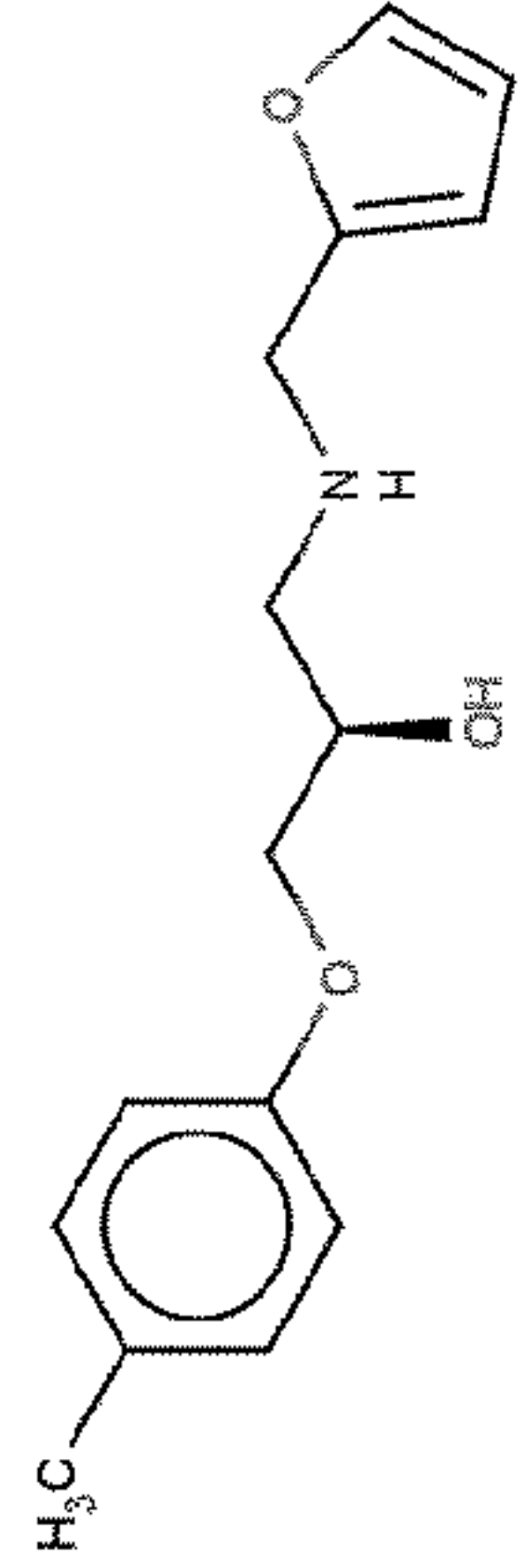
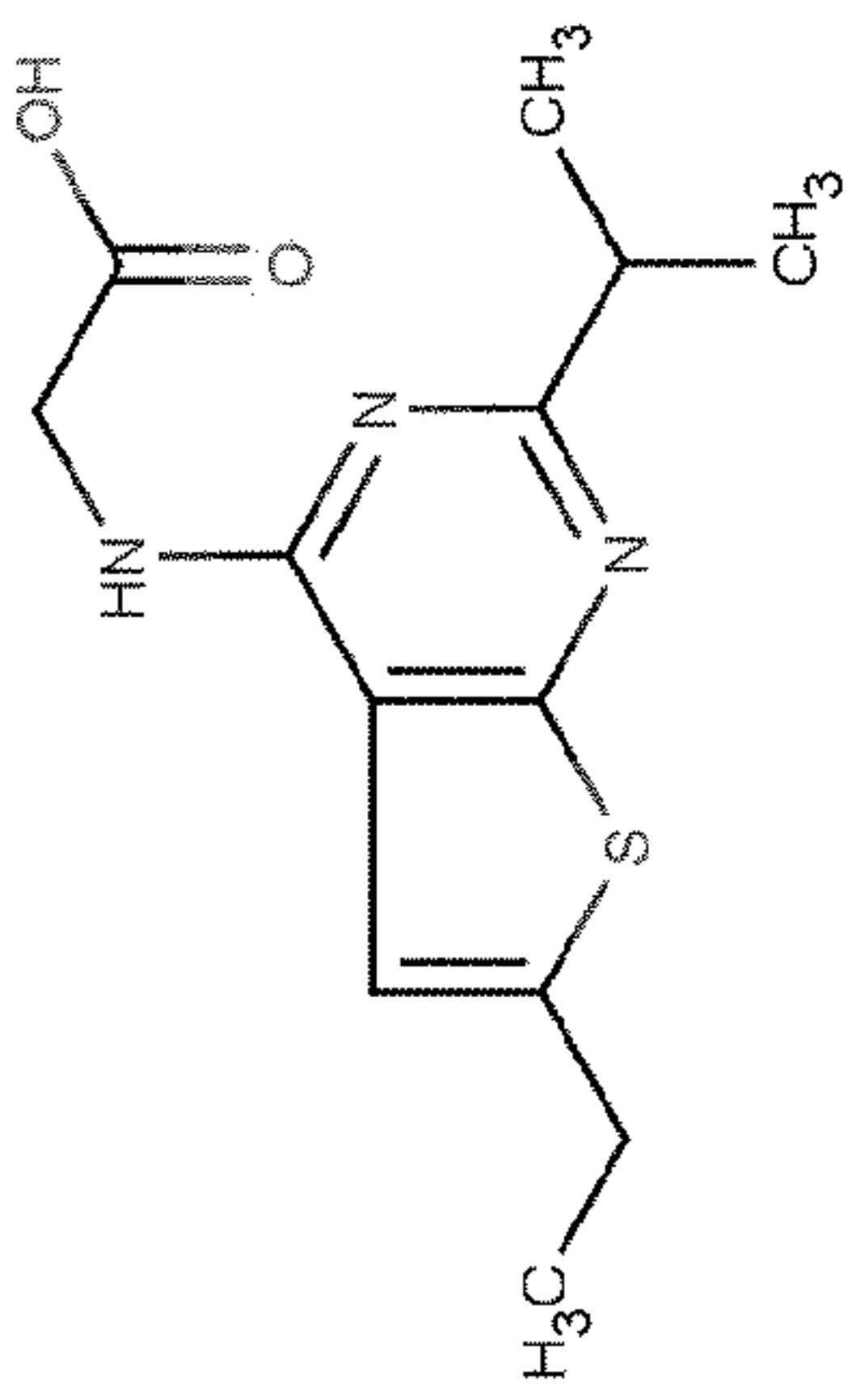
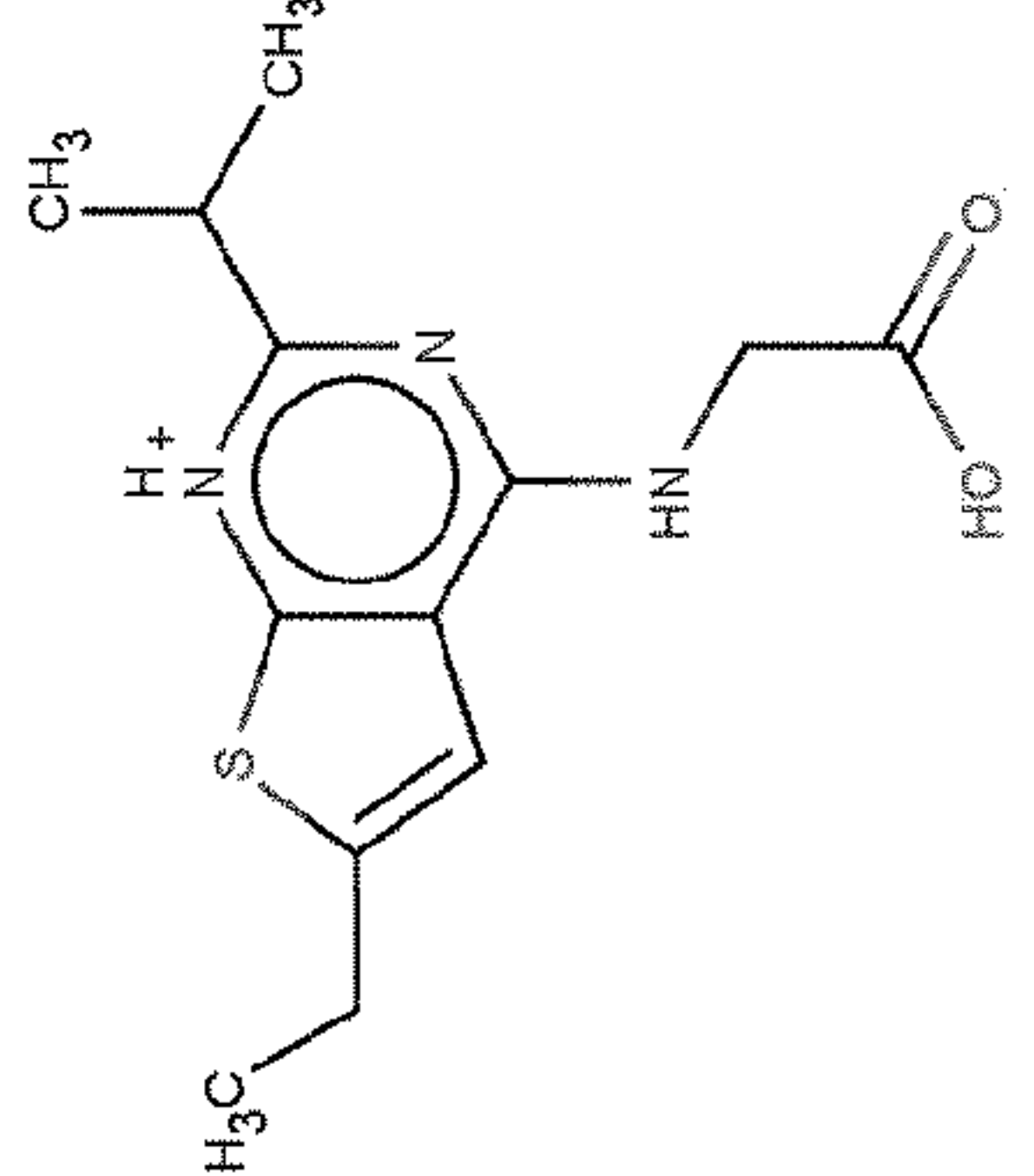
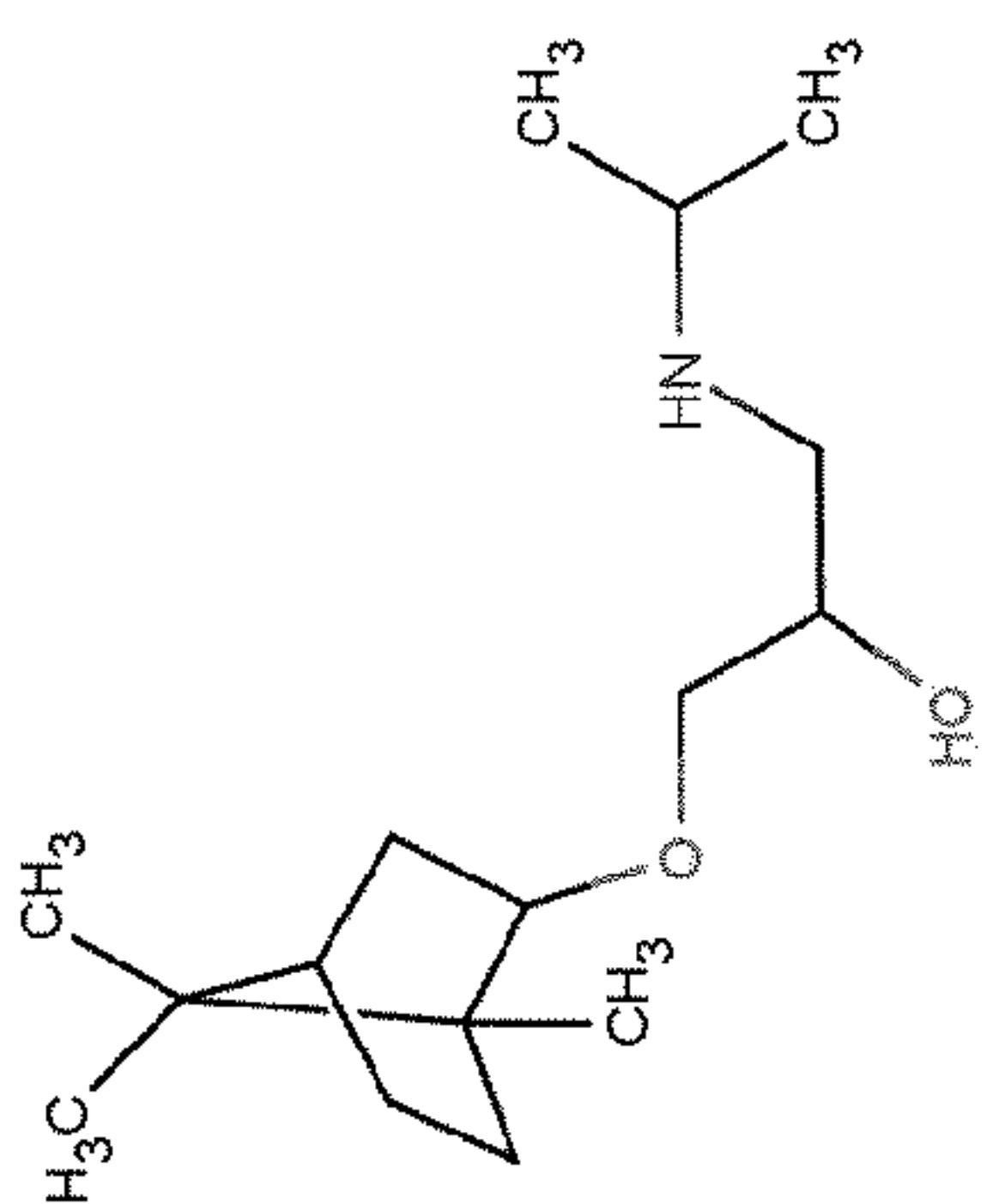
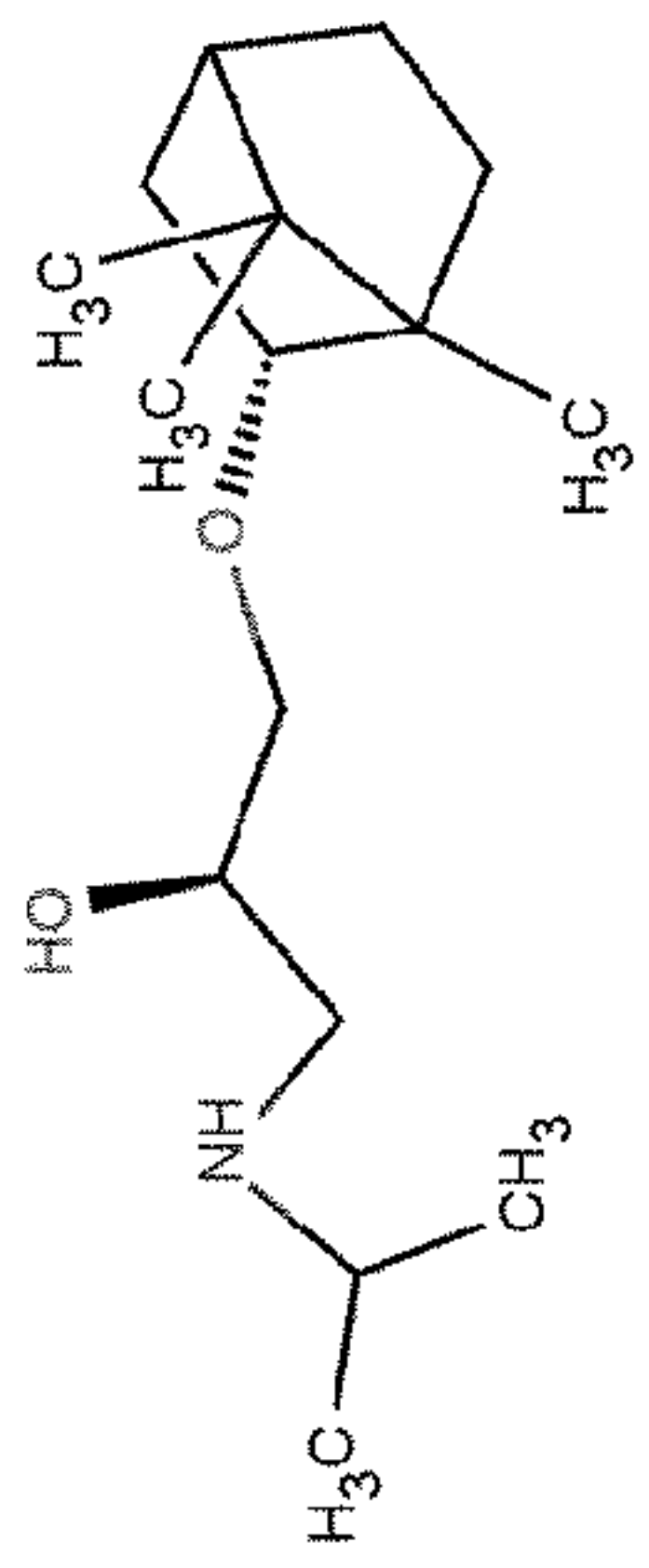
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 172		F3254-0055		F3254-0055_1_3 (1 of 8)
Compound 173		F3254-0056		F3254-0056_1 (1 of 4)
Compound 174		F3254-0059		F3254-0059_1_1 (1 of 4)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 175		F3254-0063		F3254-0063_1 (1 of 4)
Compound 176		F3254-0064		F3254-0064_1 (1 of 4)
Compound 177		F3254-0065		F3254-0065_1_1 (1 of 4)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 178		F3254-0073		F3254-0073_1_1 (1 of 8)
Compound 179		F3254-0075		F3254-0075_1 (1 of 4)
Compound 180		F3254-0098		F3254-0098_1_2 (1 of 4)

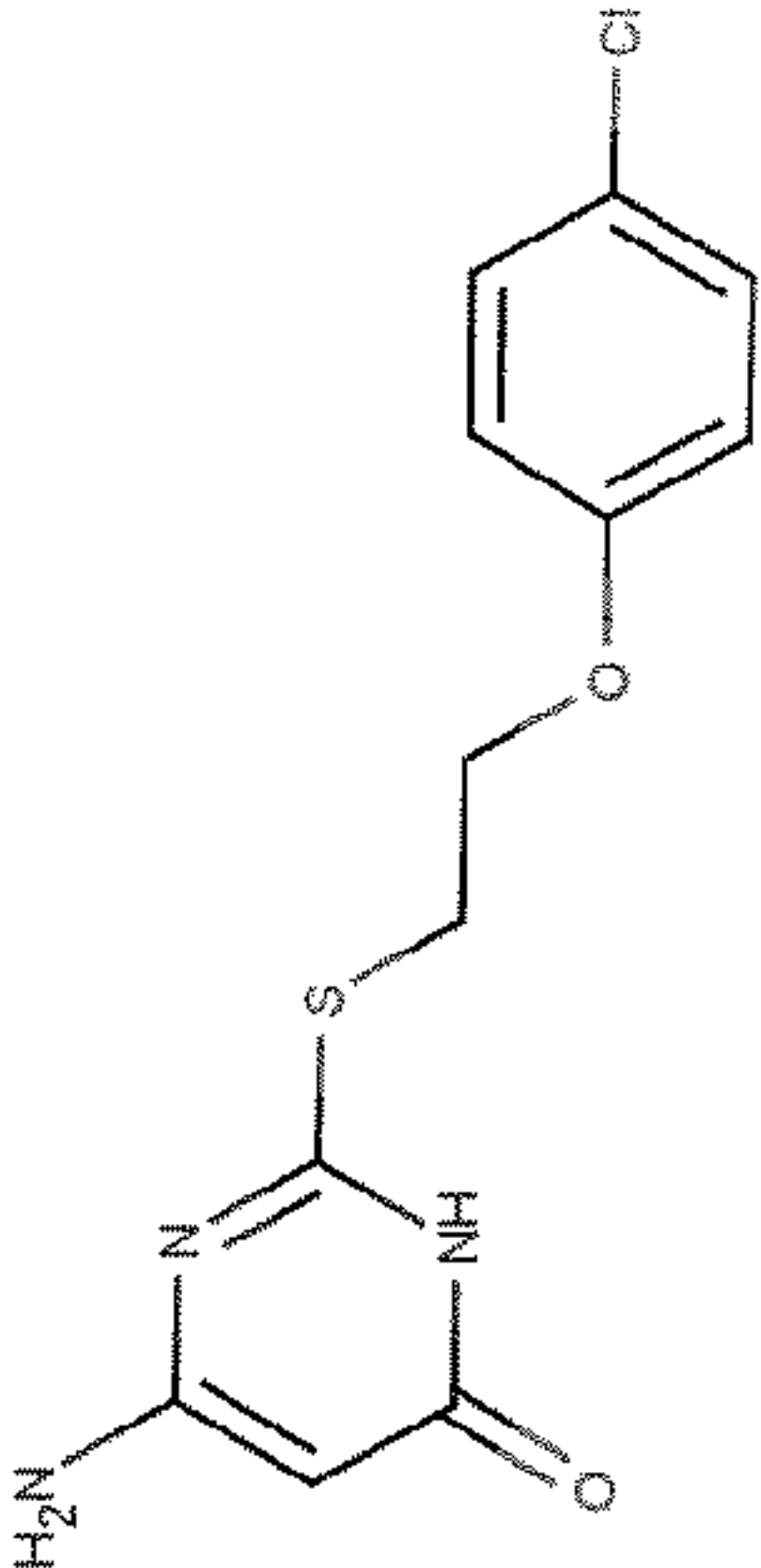
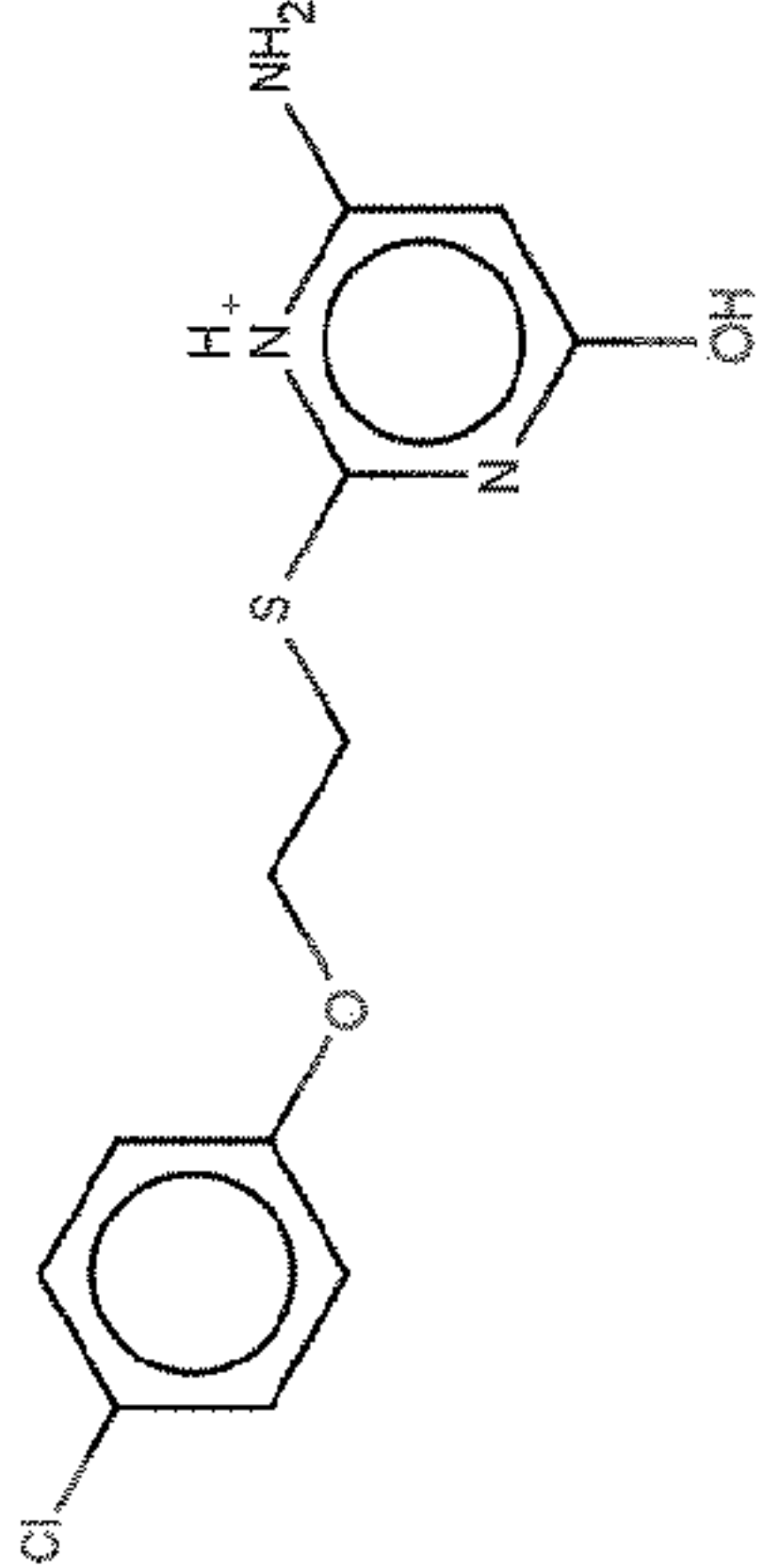
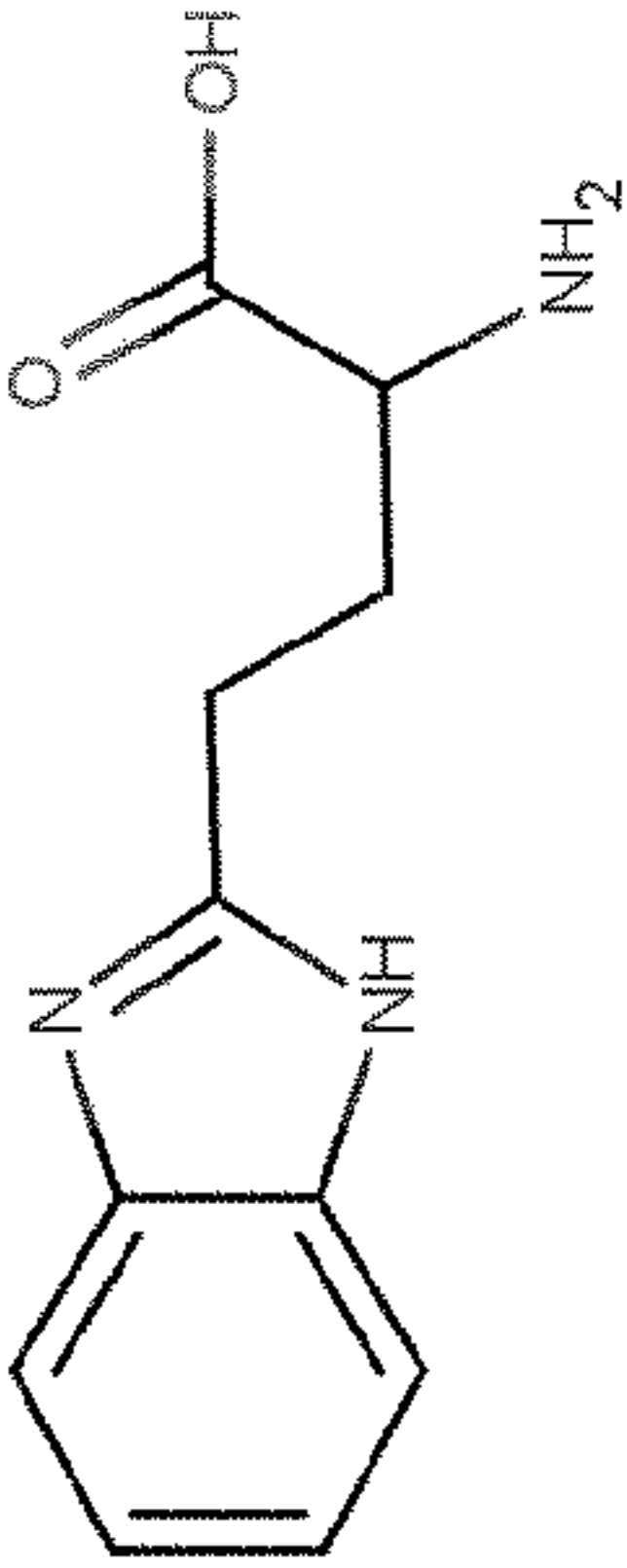
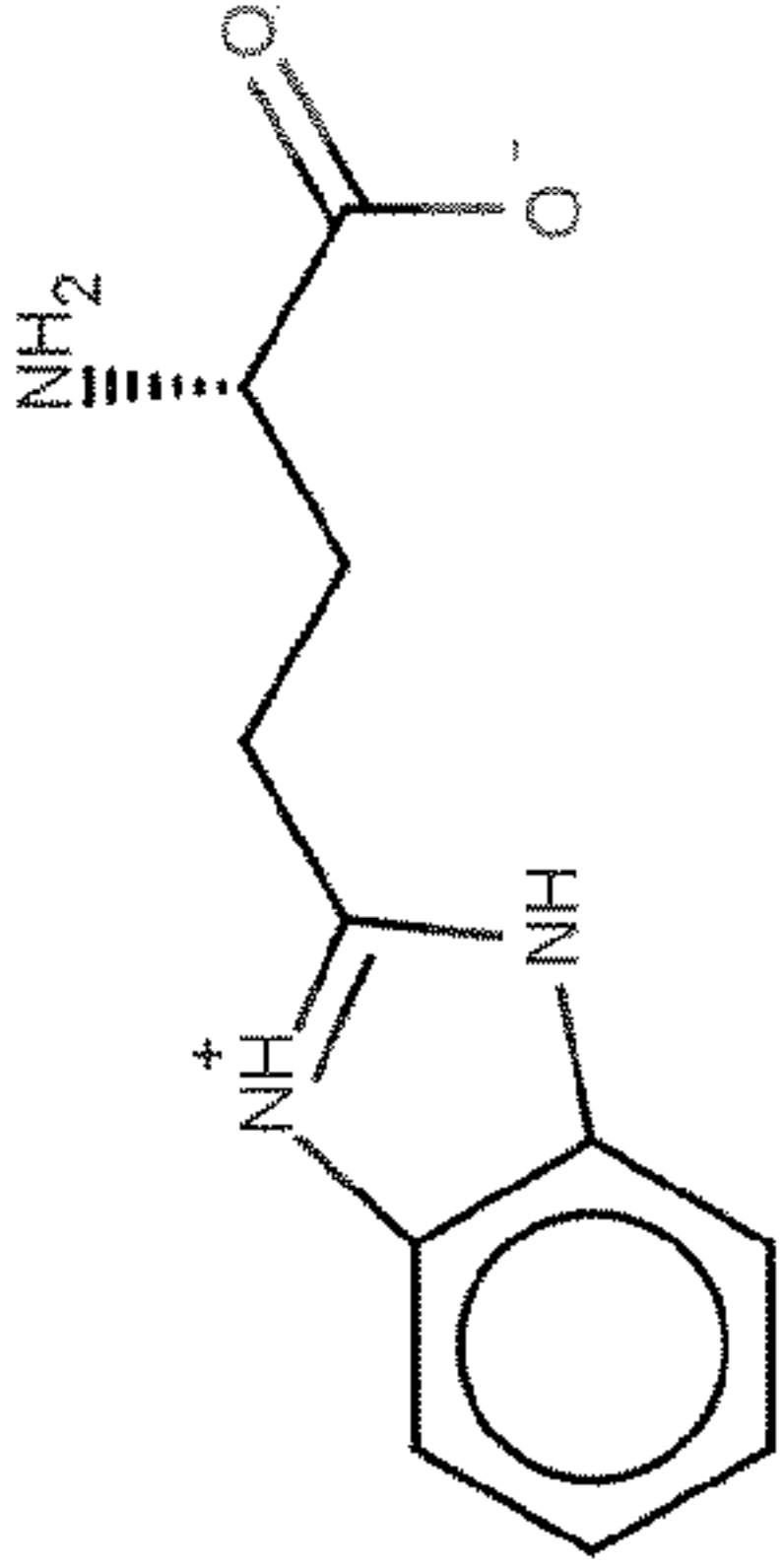
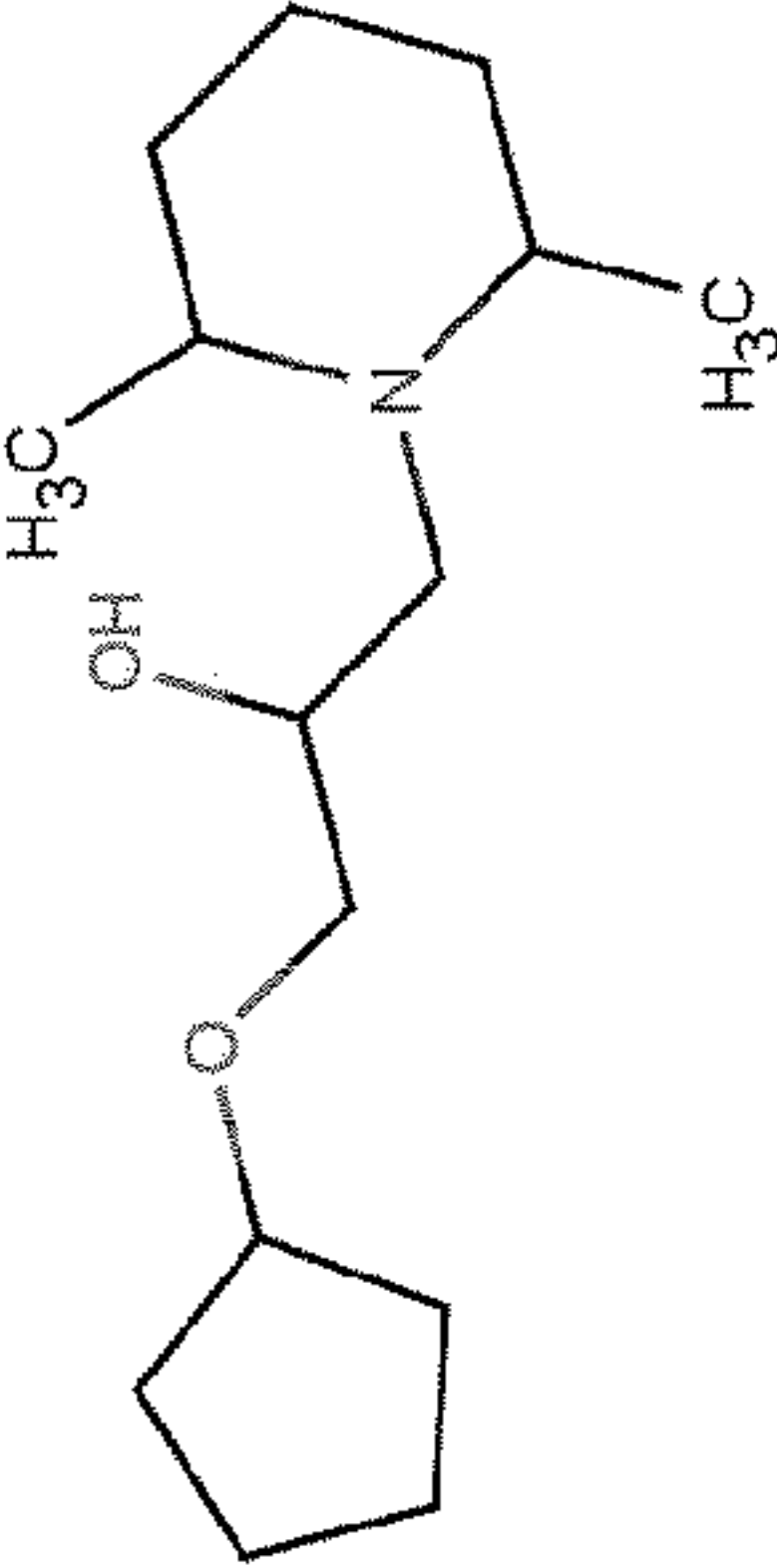
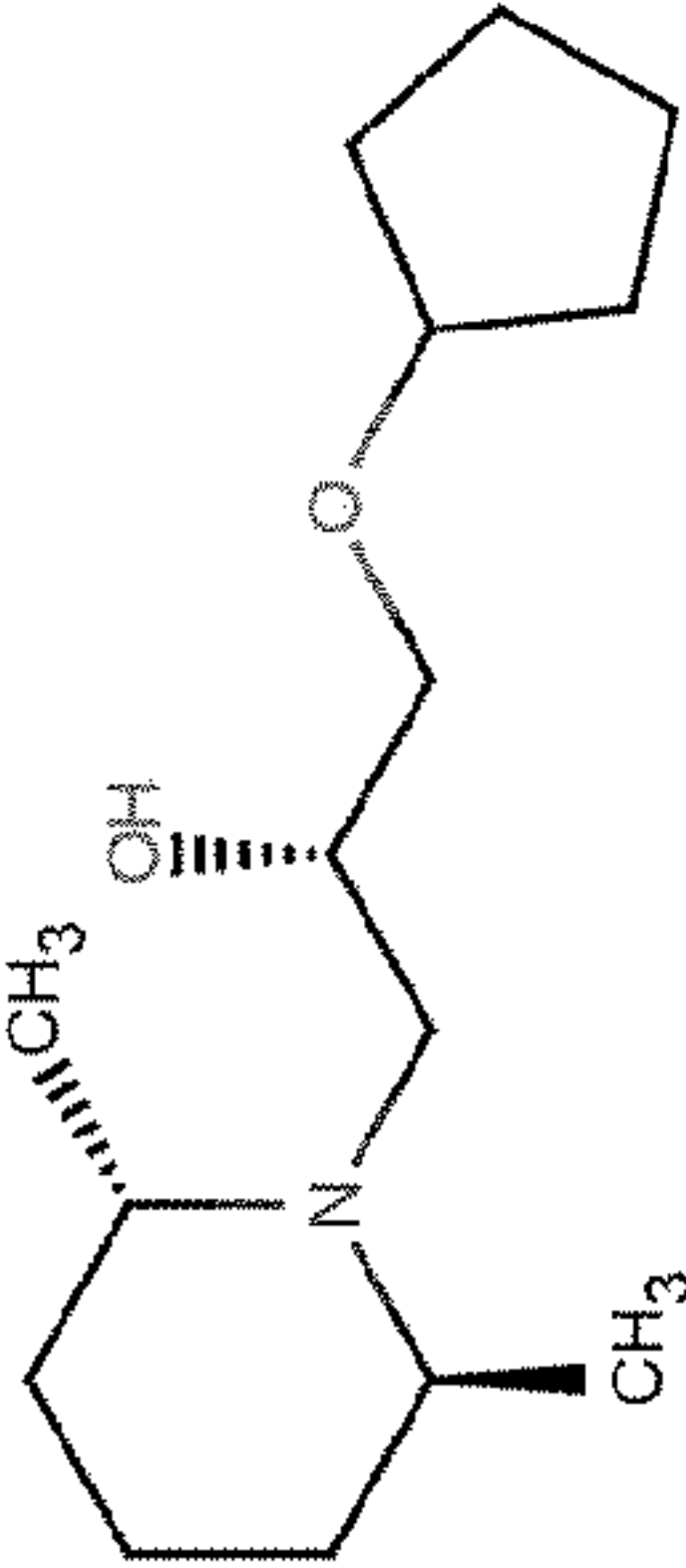
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 181		F3254-0105		F3254-0105_1_6 (1 of 16)
Compound 182		F3255-0157		F3255-0157_1_1 (1 of 8)
Compound 183		F3260-0881		F3260-0881_7_1 (1 of 13)

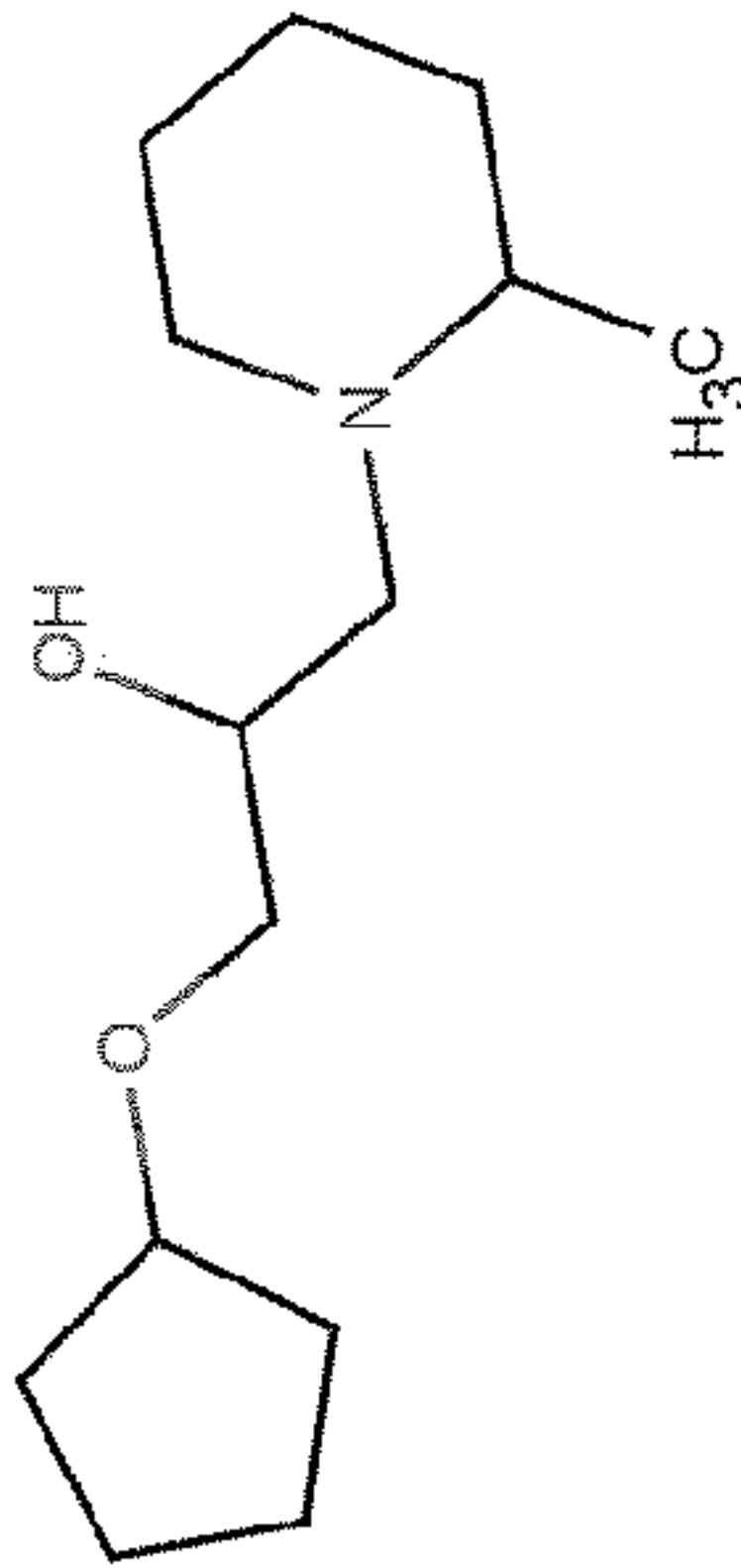
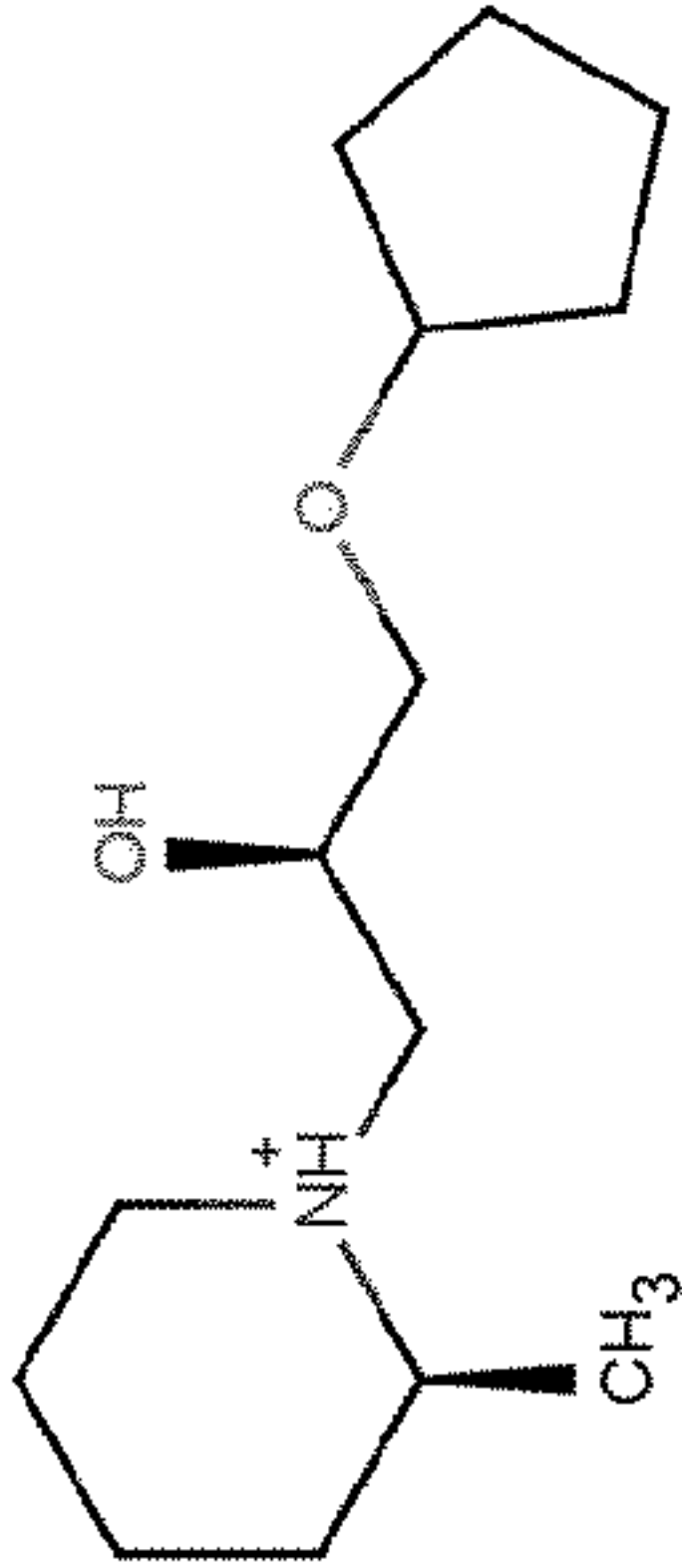
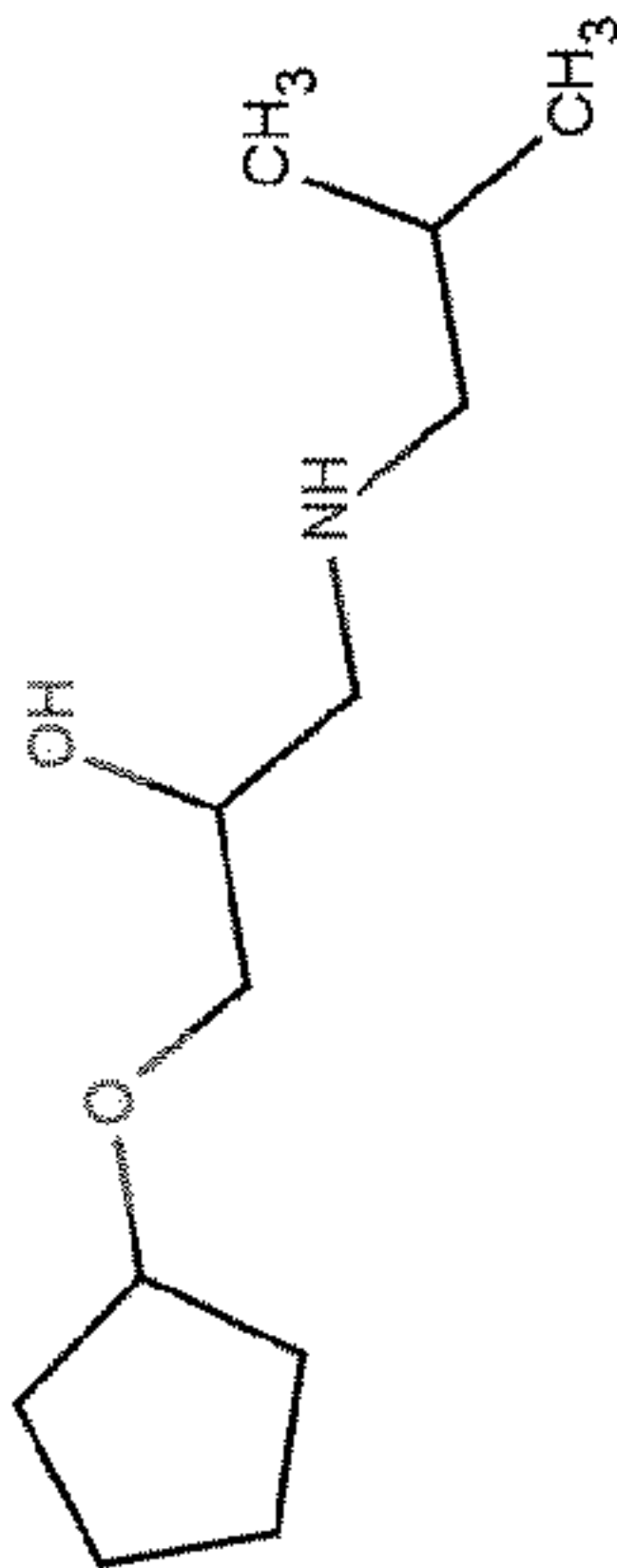
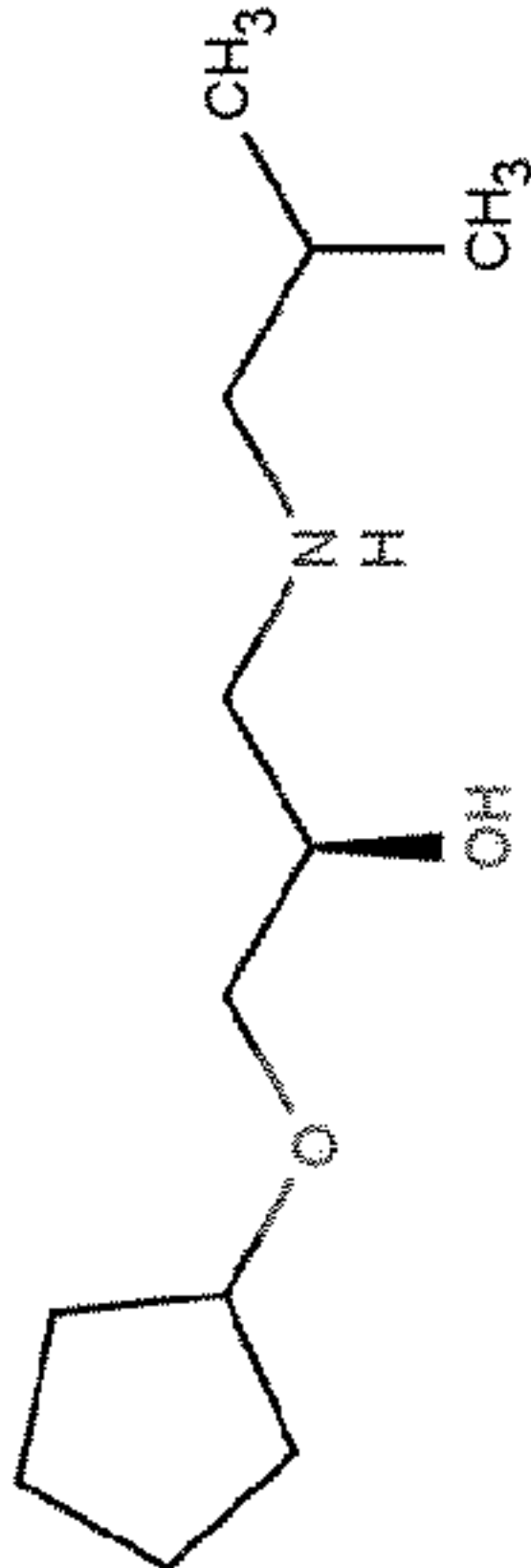
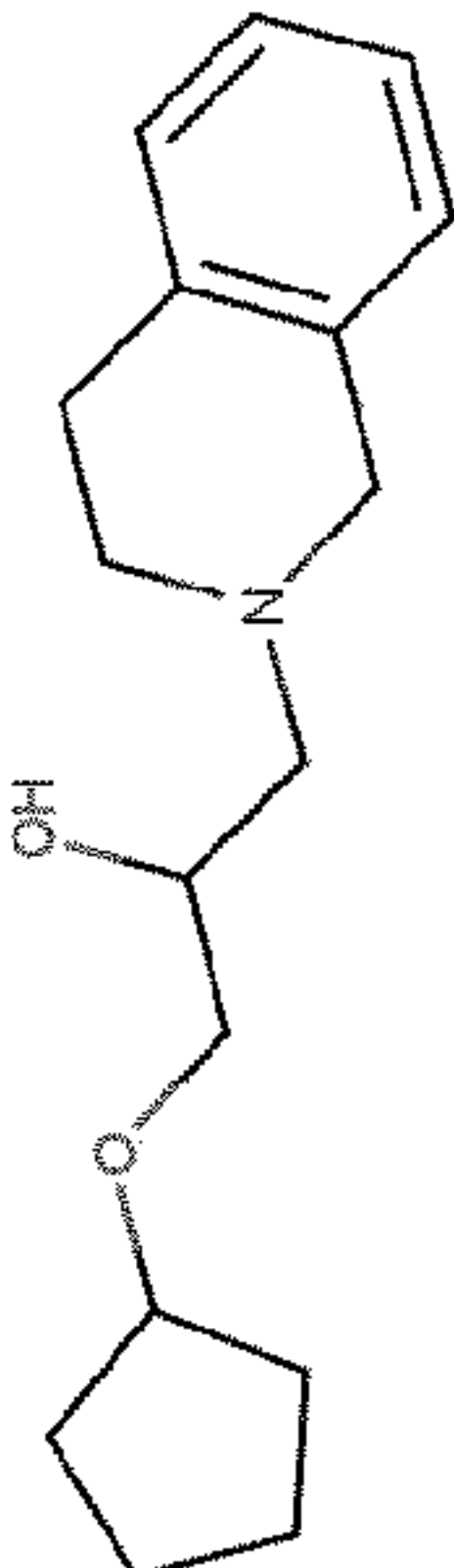
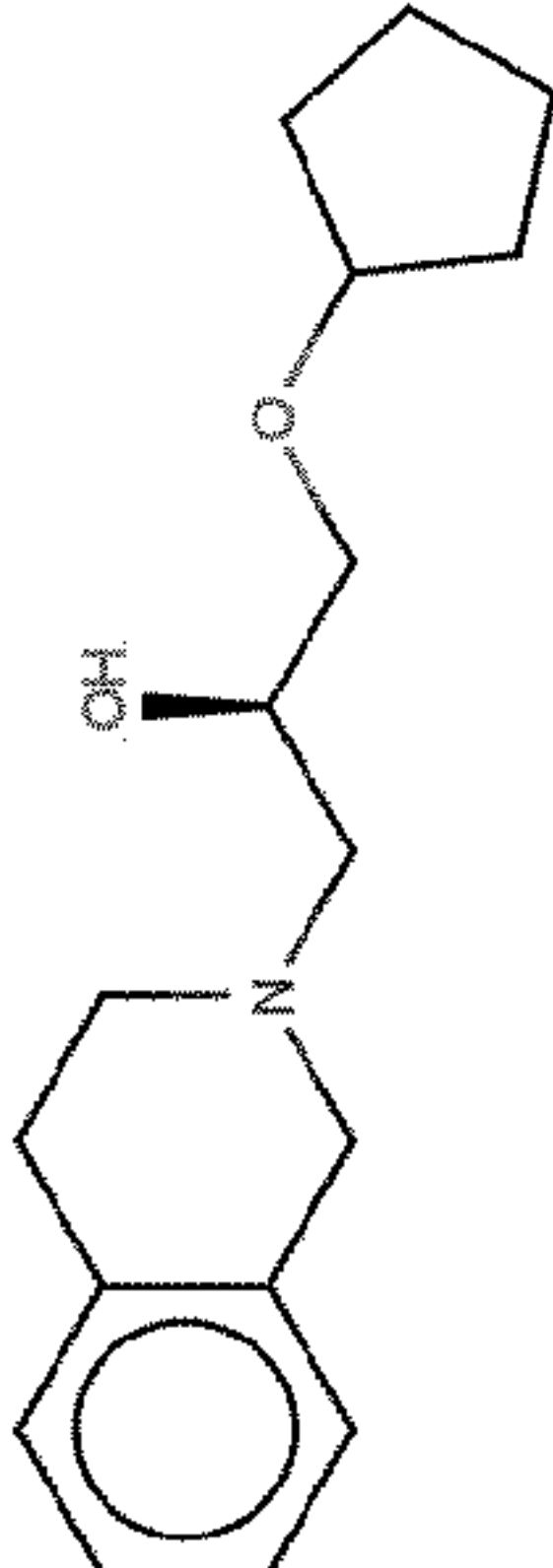
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 184		F3266-0086		F3266-0086_1_1 (1 of 2)
Compound 185		F3276-0018		F3276-0018_2 (1 of 4)
Compound 186		F3276-0031		F3276-0031_2 (1 of 4)

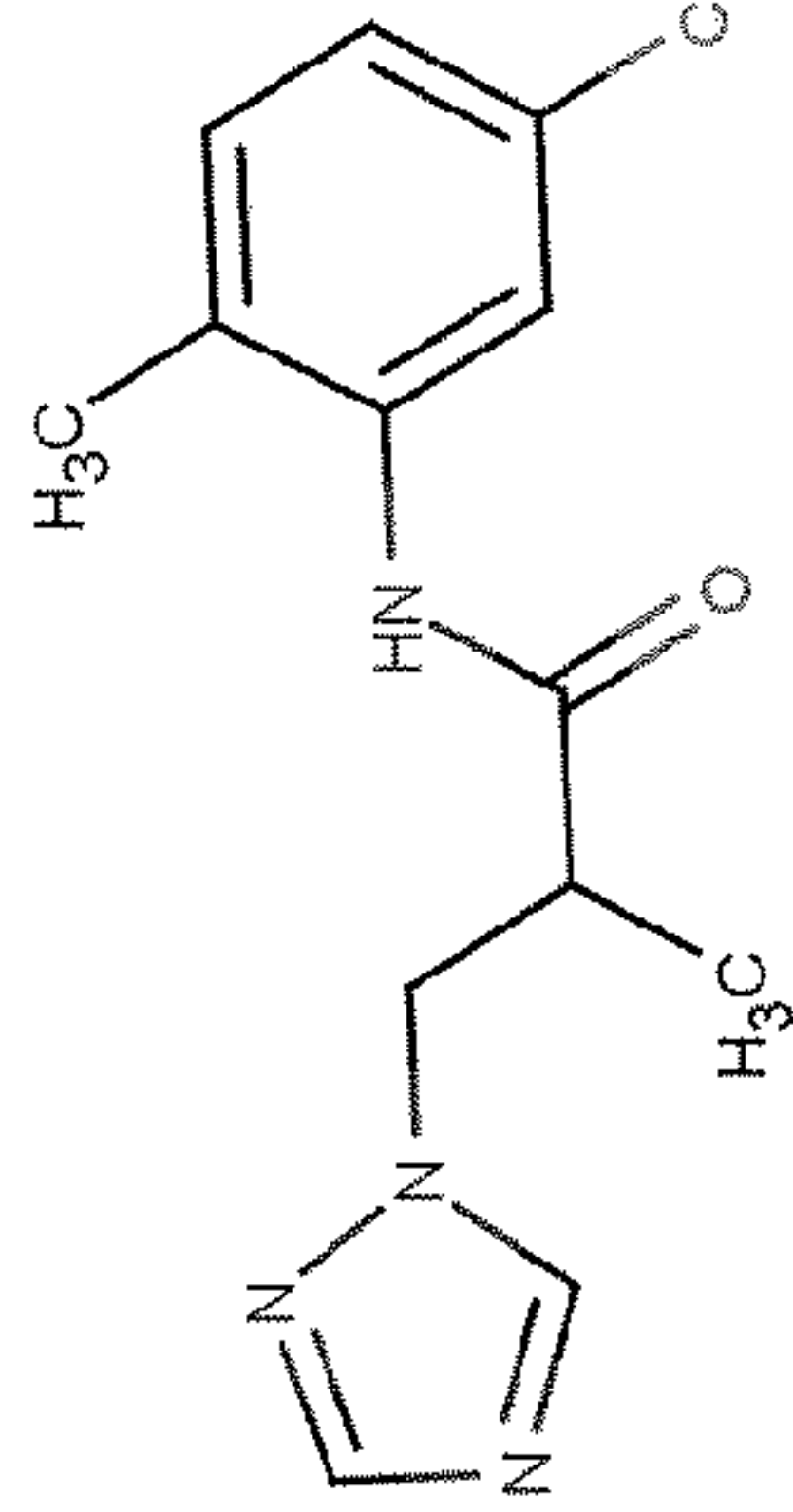
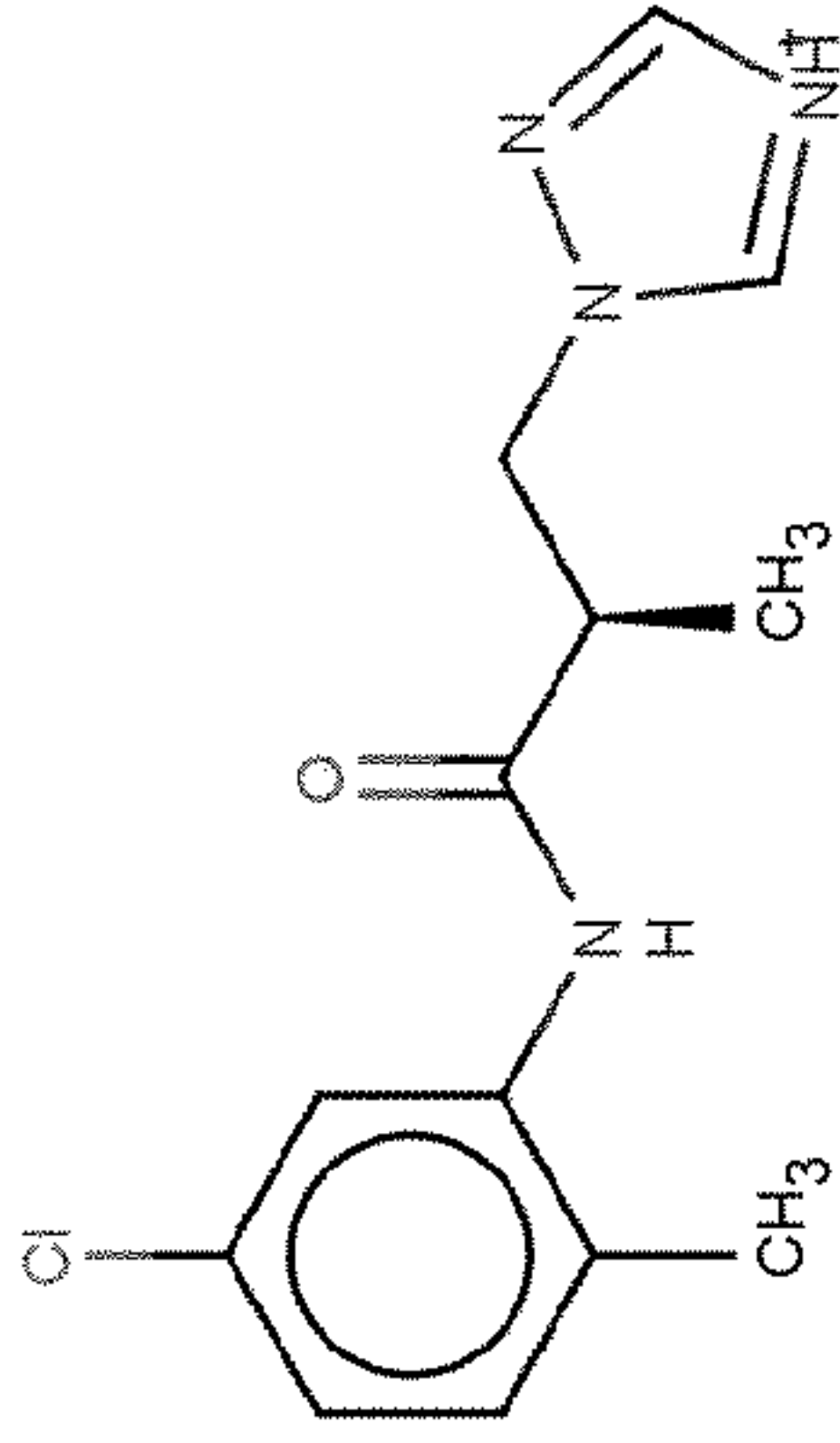
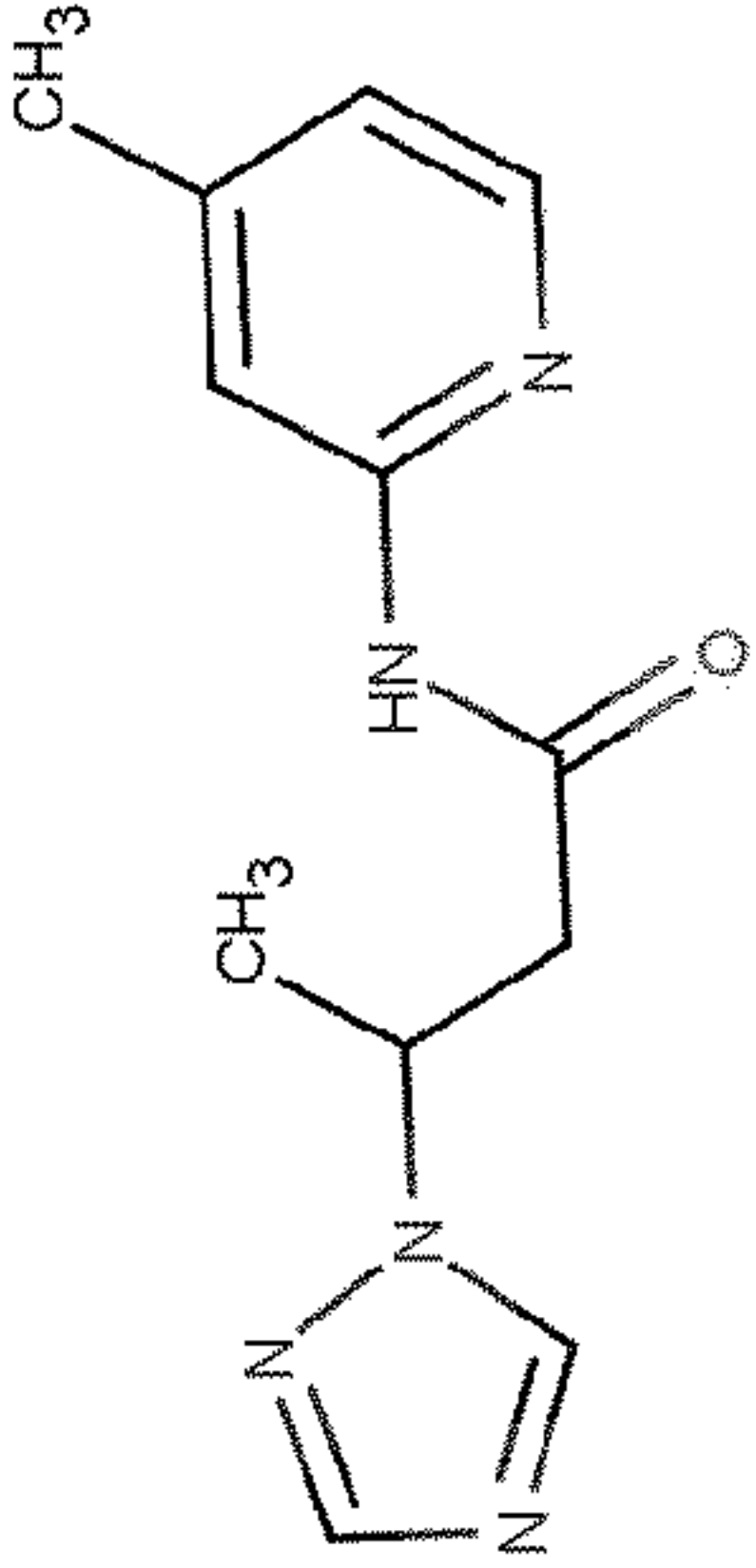
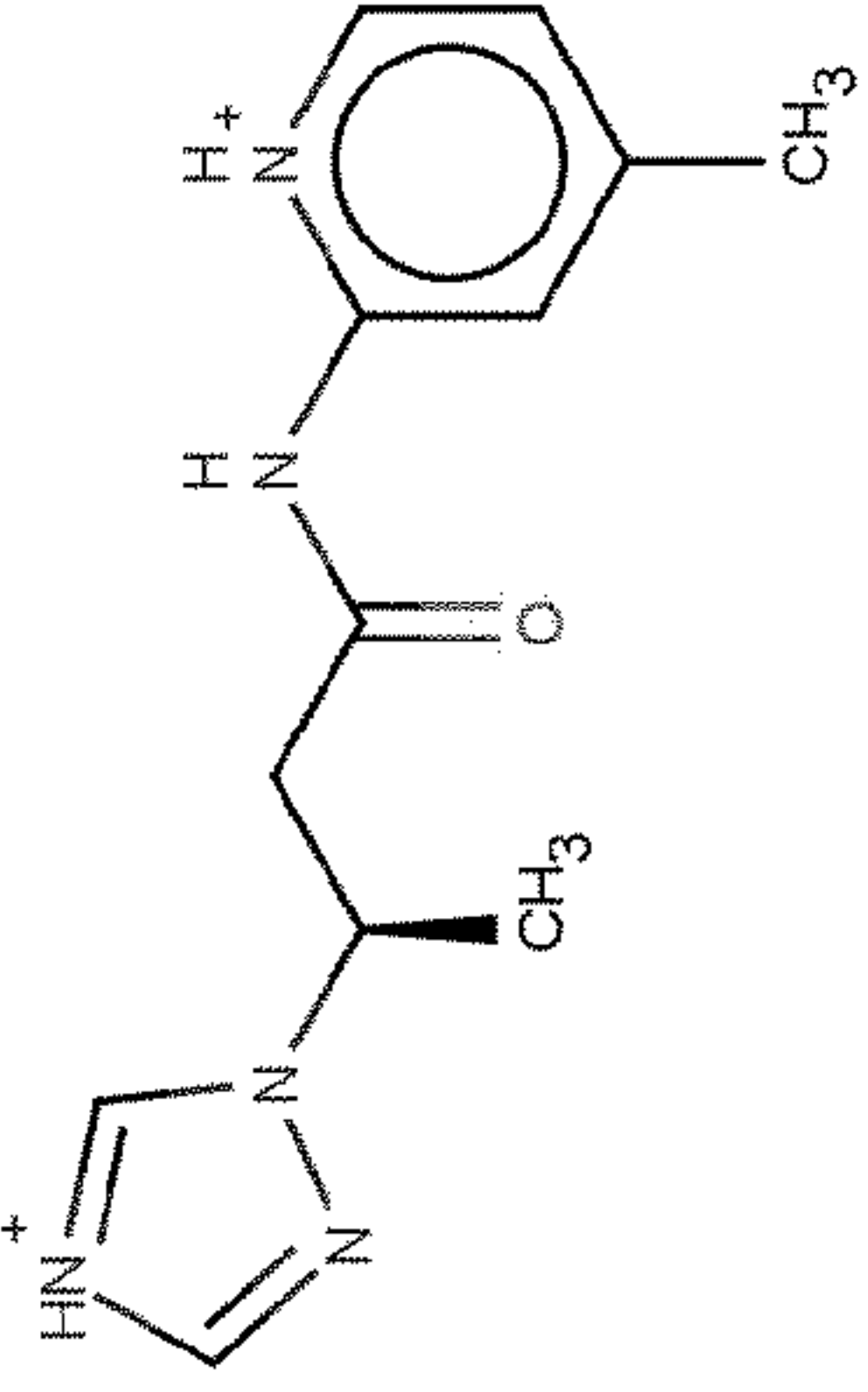
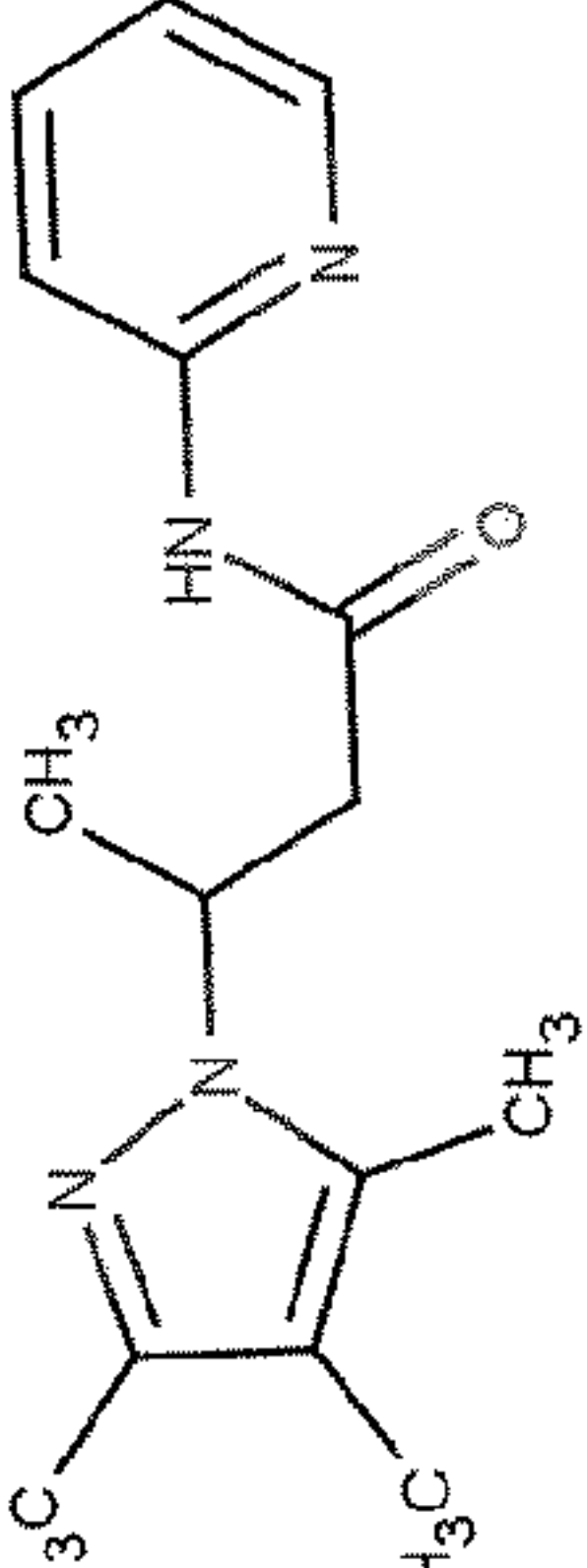
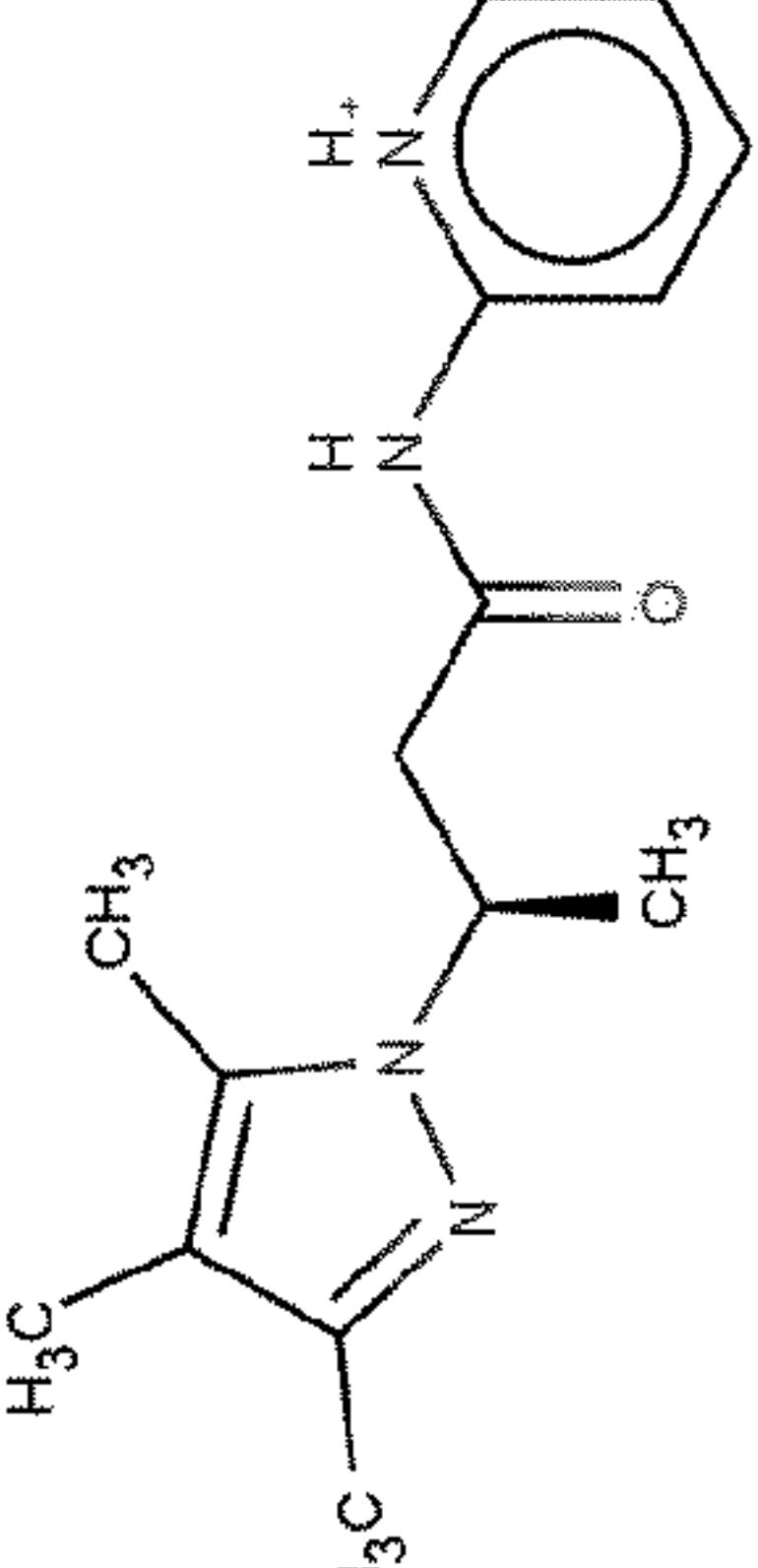
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 187		F3276-0044		F3276-0044_2 (1 of 4)
Compound 188		F3277-0798		F3277-0798_1_1 (1 of 2)
Compound 189		F3278-0013		F3278-0013_4 (1 of 8)

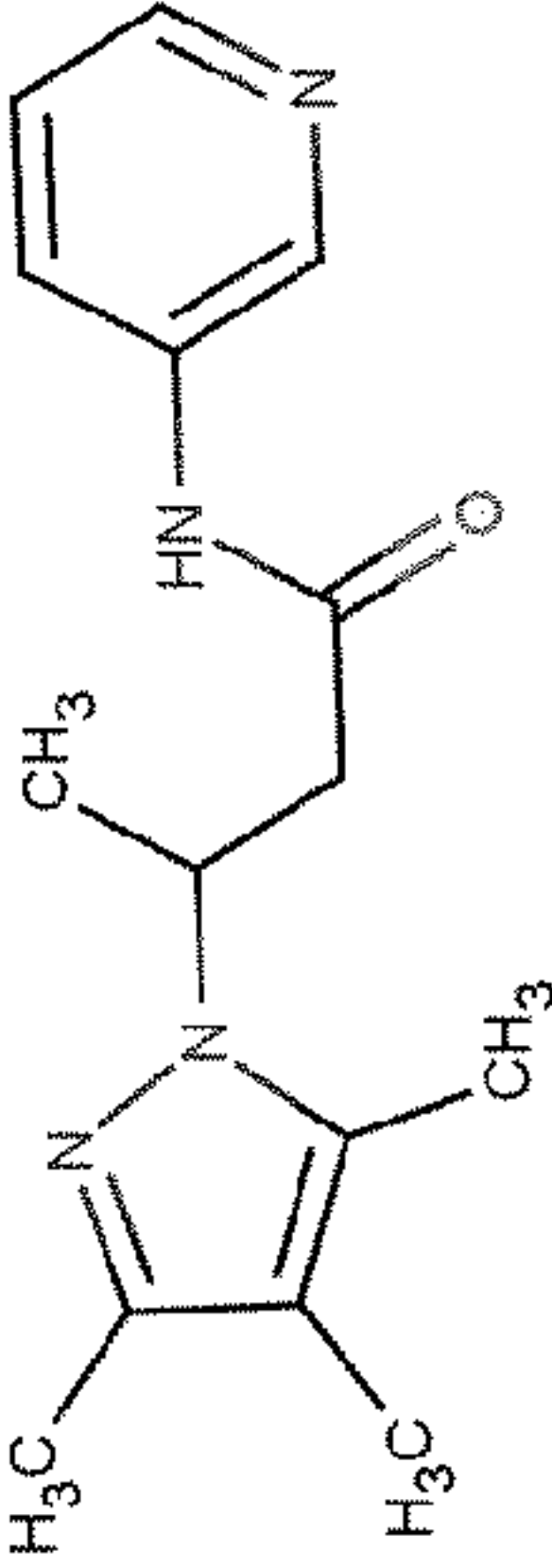
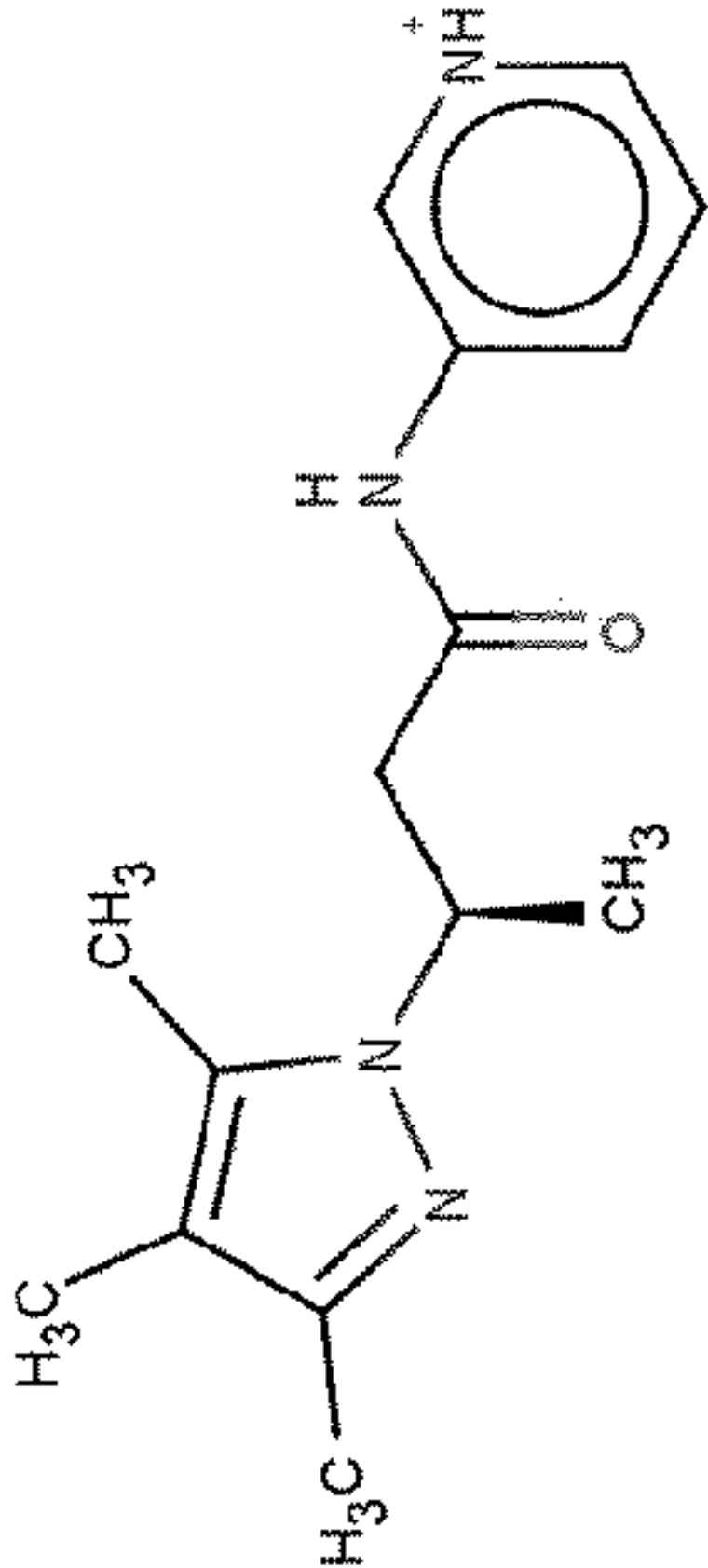
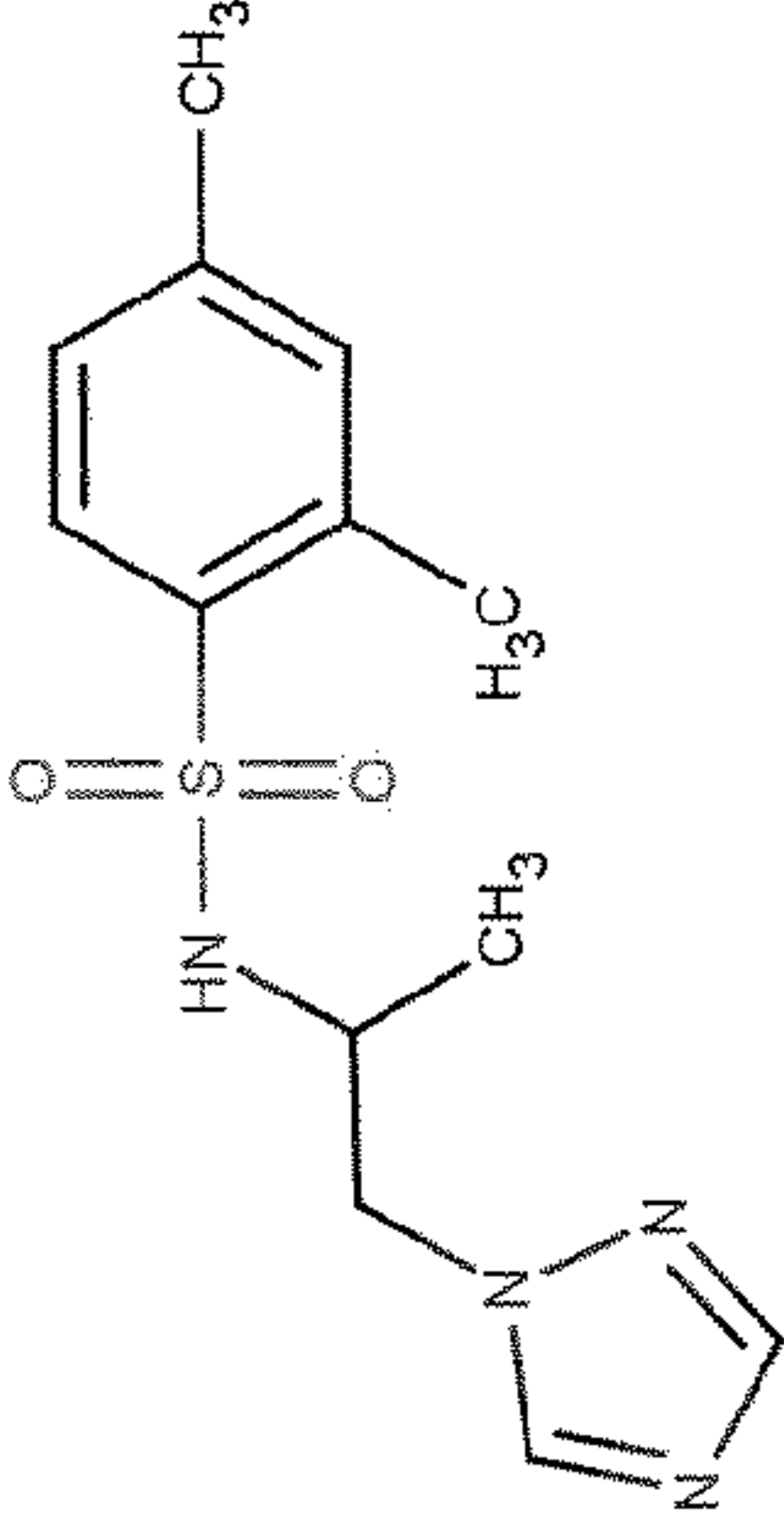
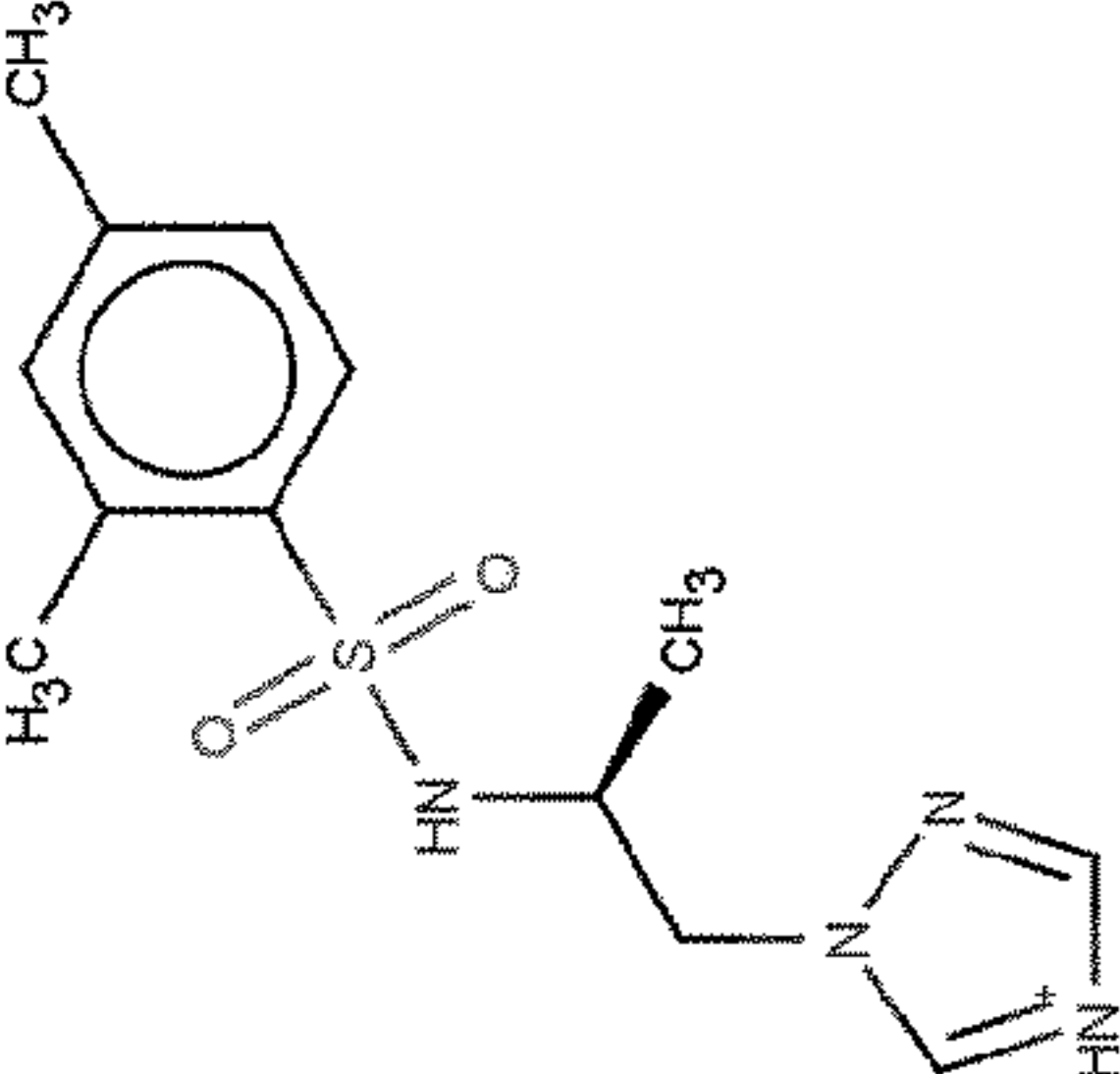
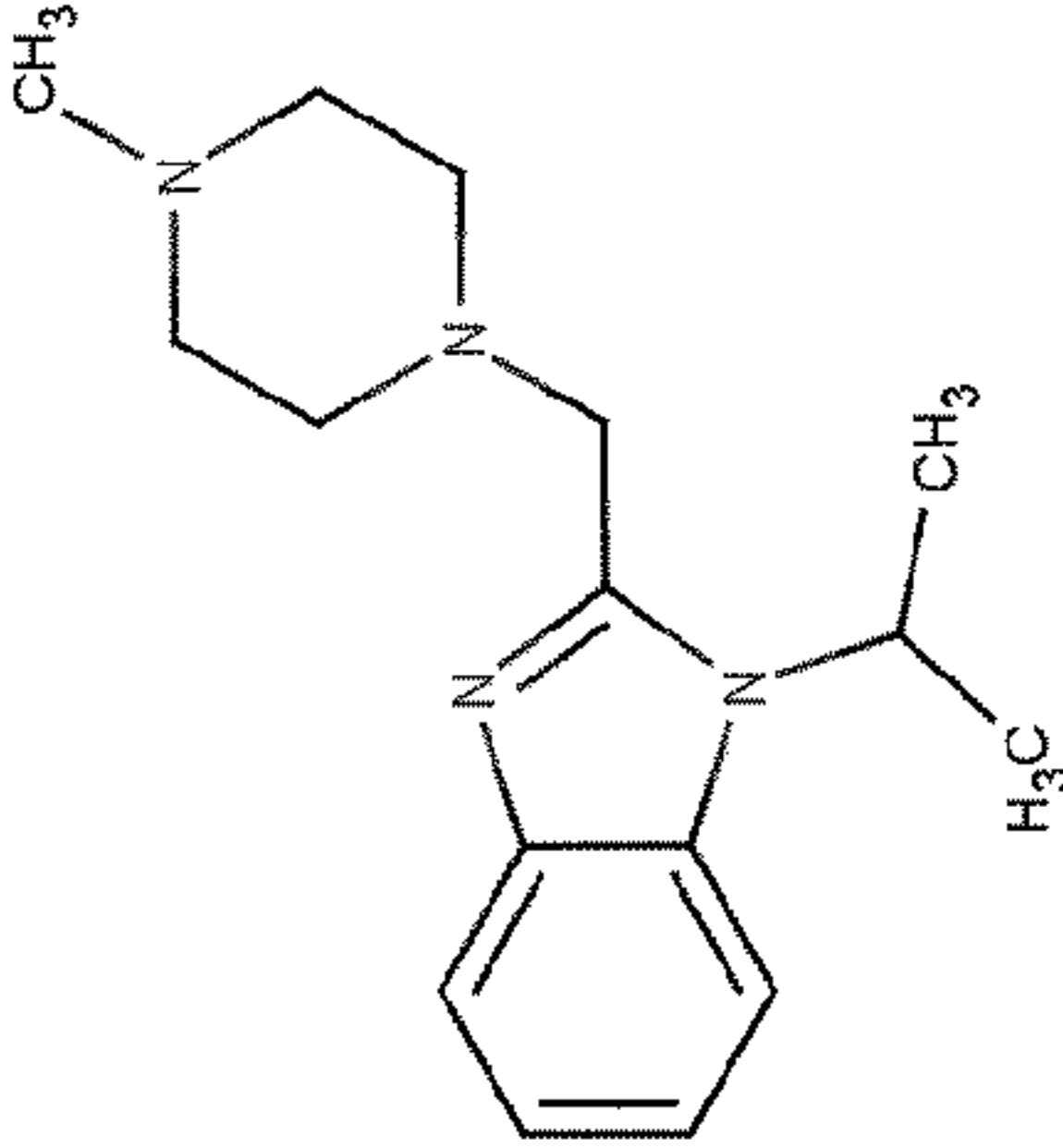
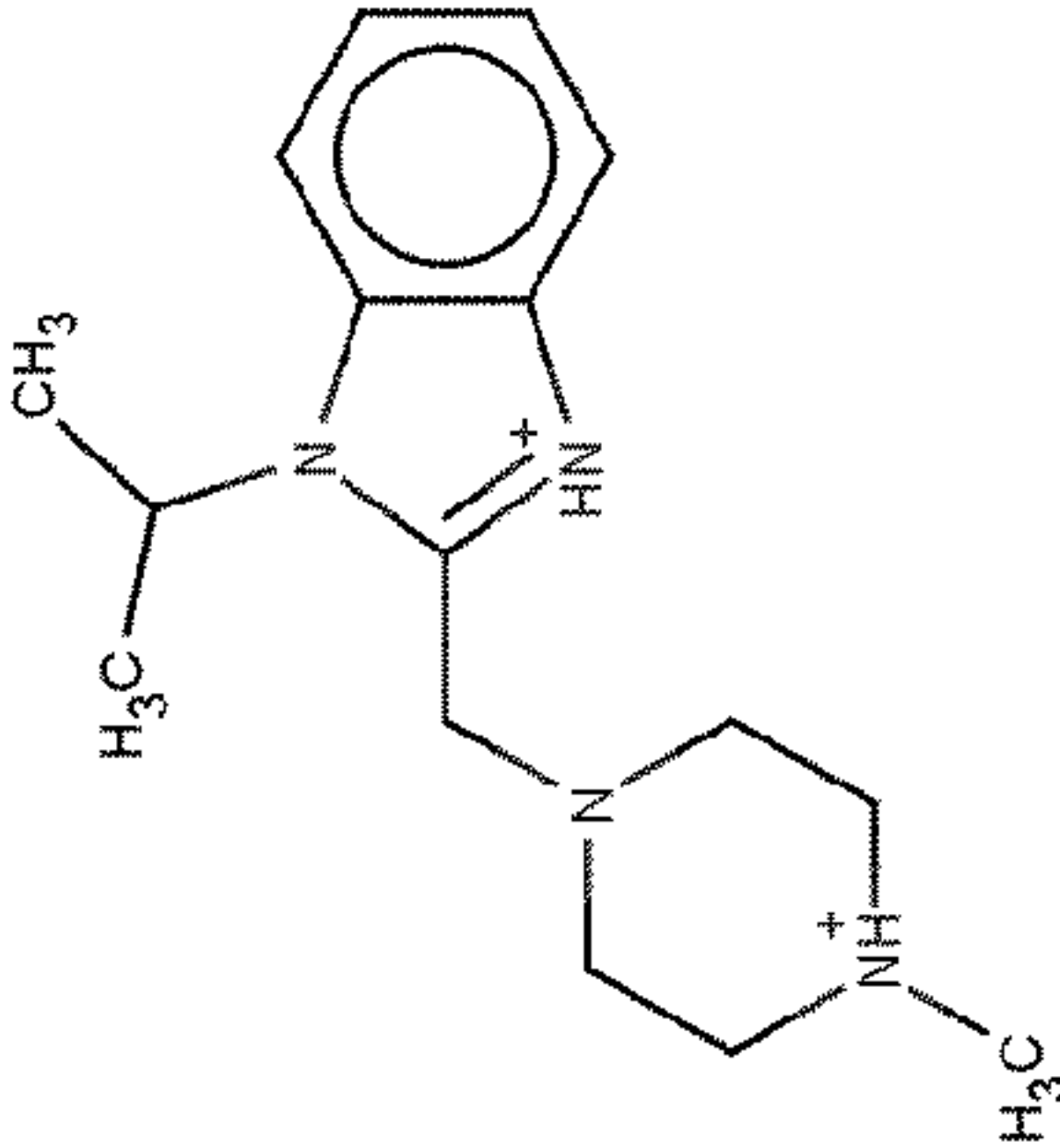
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 190		F3284-7619		F3284-7619_2 (1 of 20)
Compound 191		F3284-7869		F3284-7869_1_1 (1 of 2)
Compound 192		F3301-0129		F3301-0129_1_2 (1 of 4)

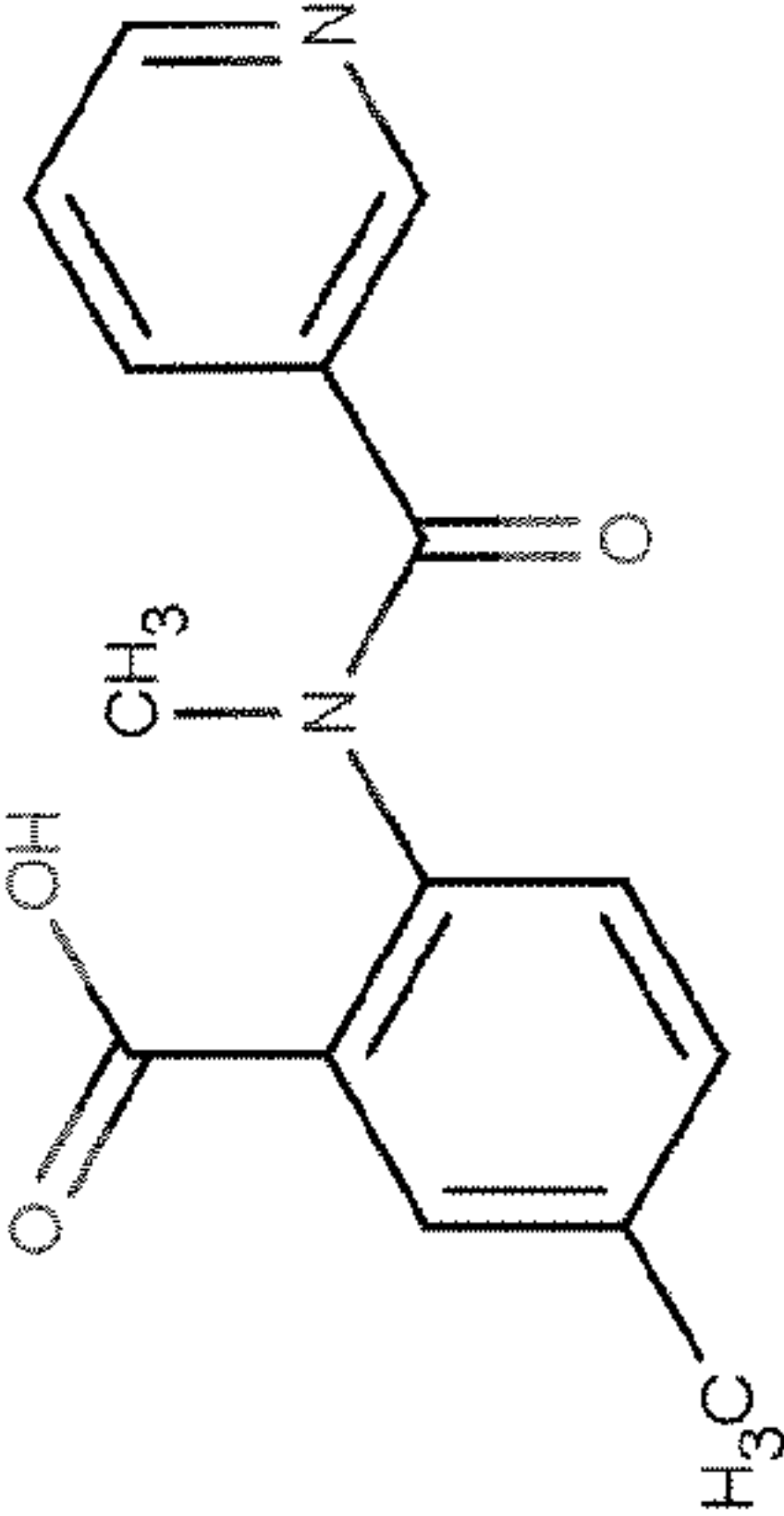
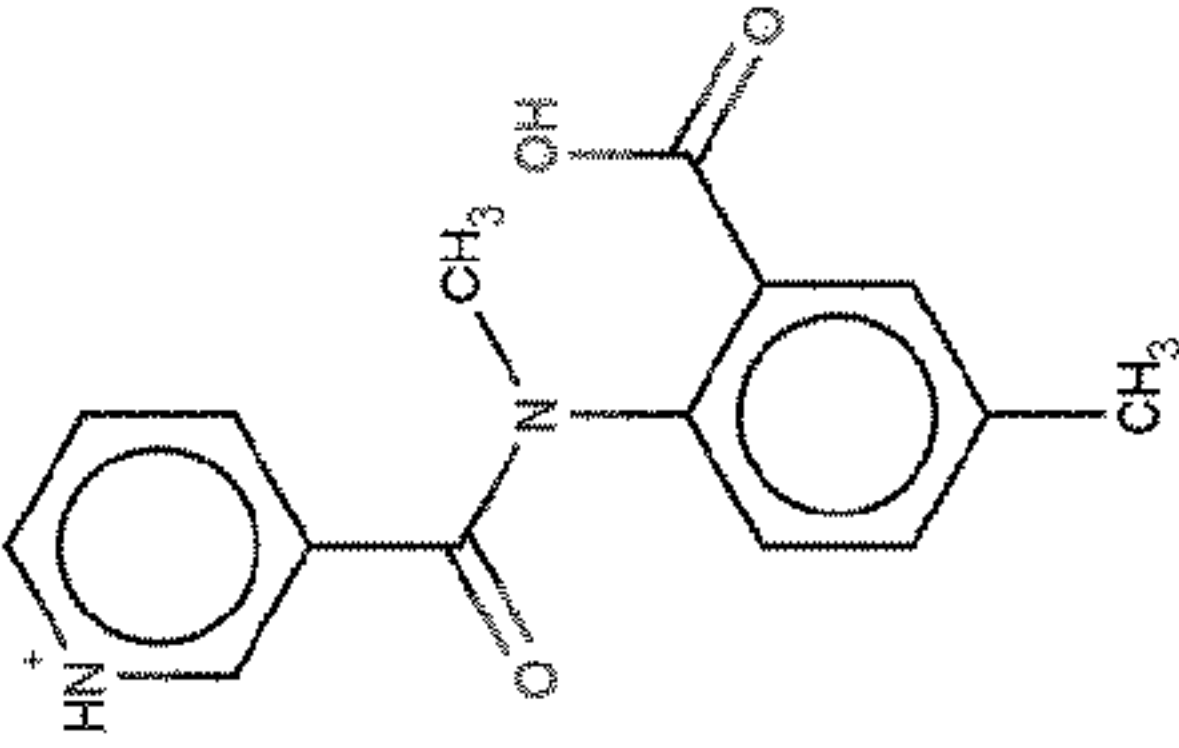
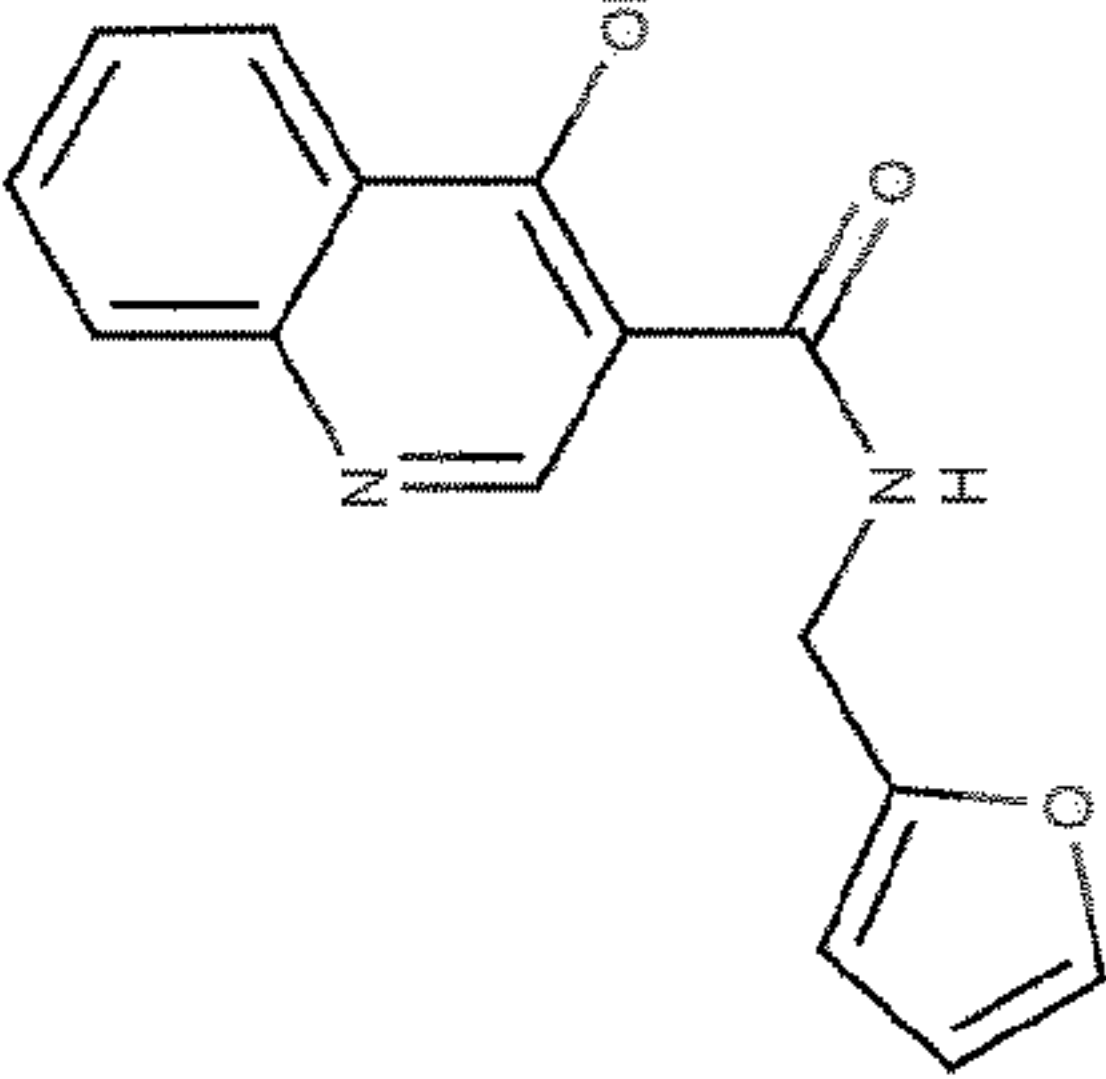
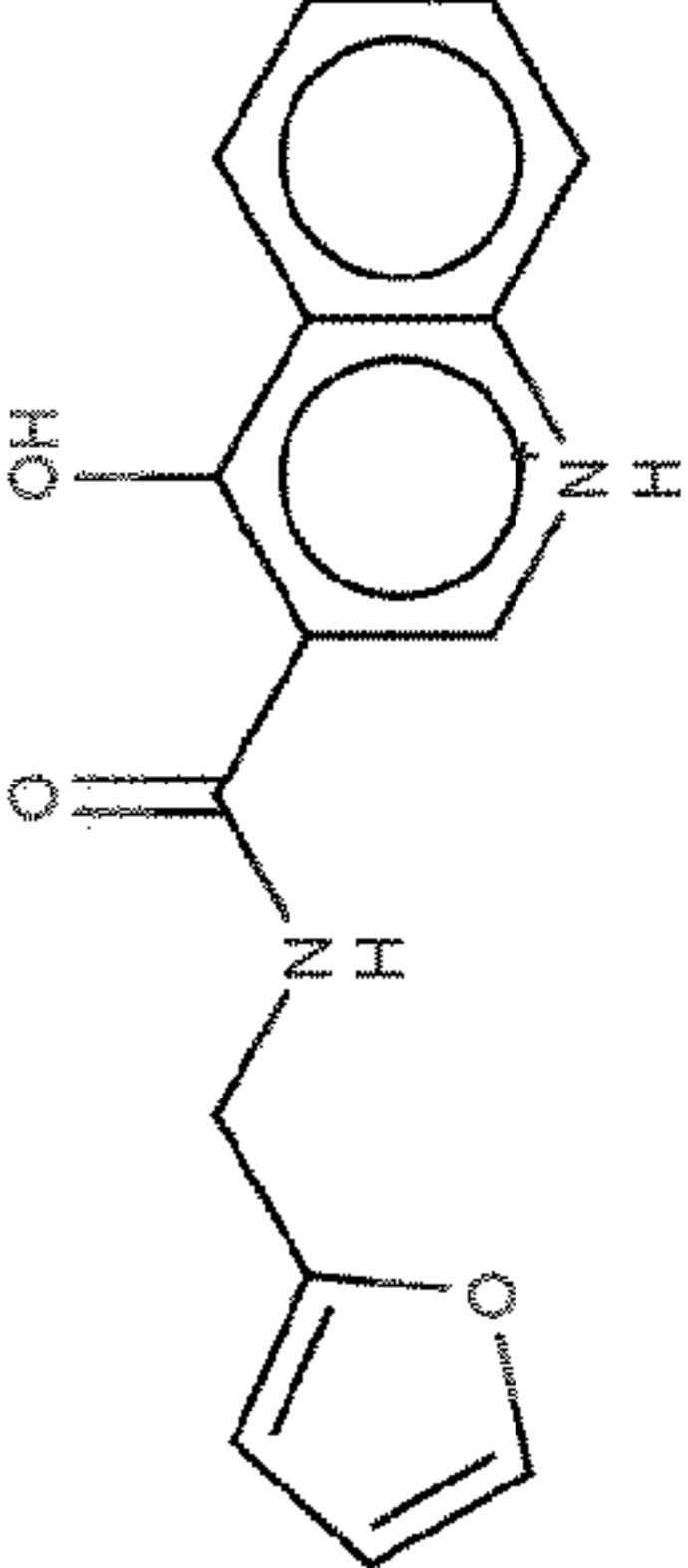
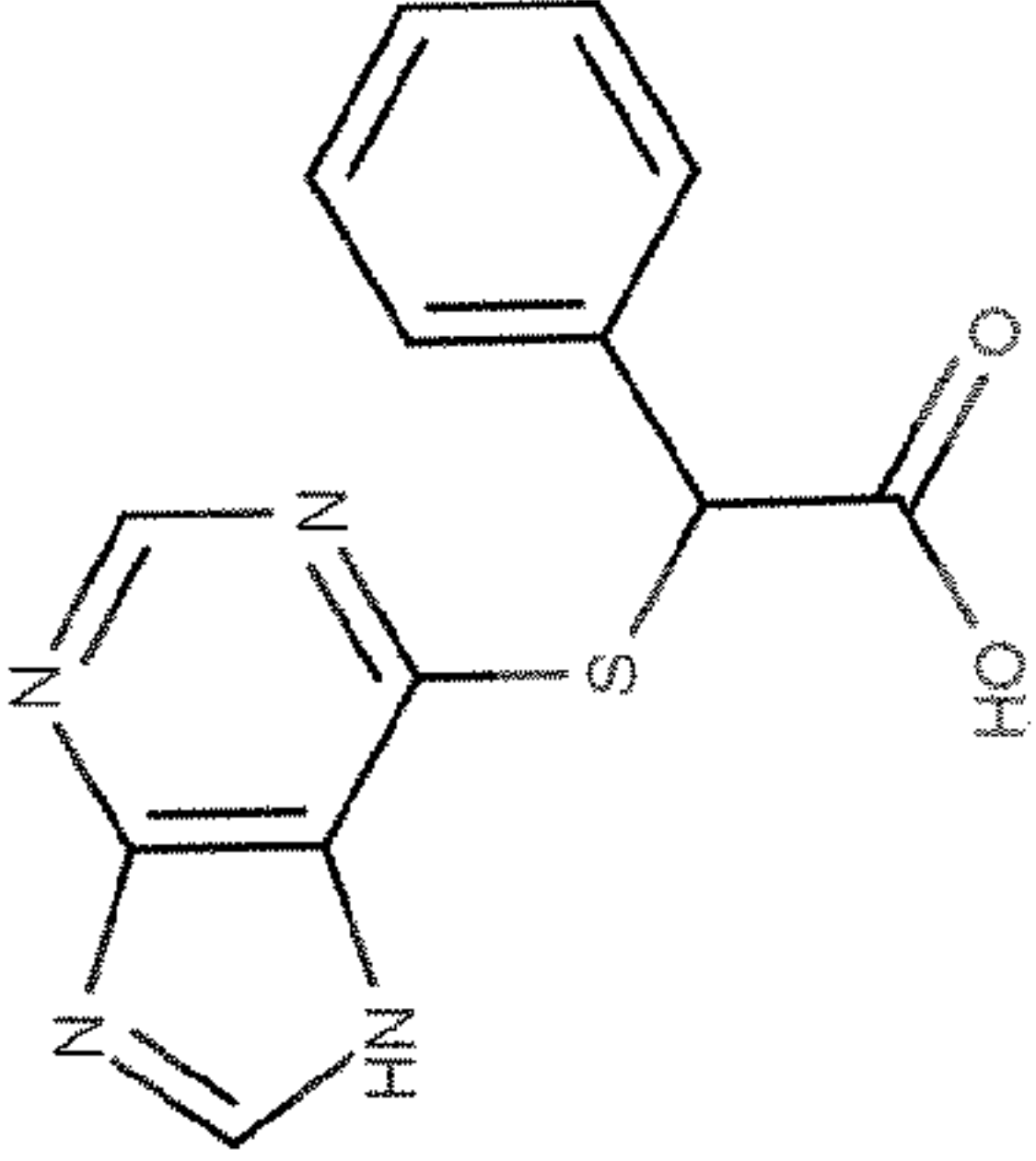
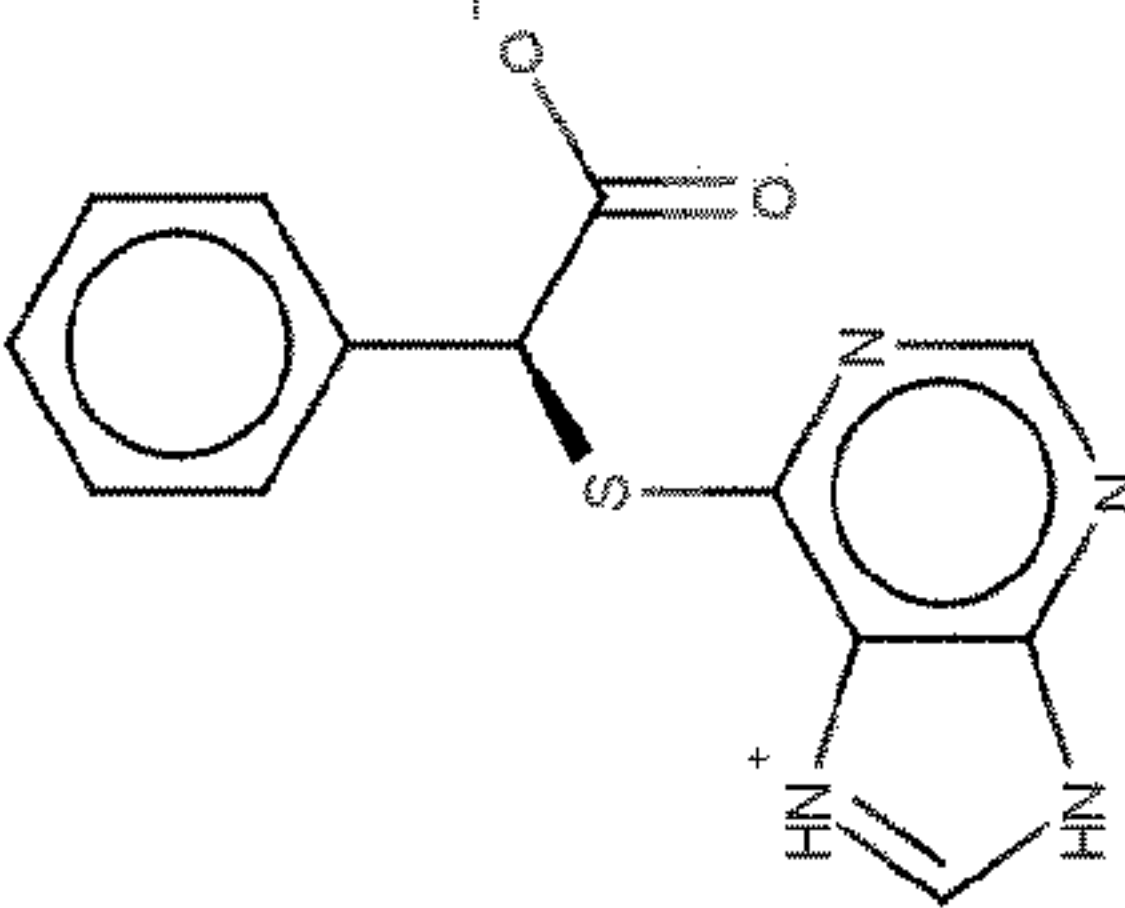
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 193		F3308-0605		F3308-0605_7_2 (1 of 16)
Compound 194		F3308-1206		F3308-1206_1 (1 of 8)
Compound 195		F3308-2774		F3308-2774_5_1 (1 of 14)

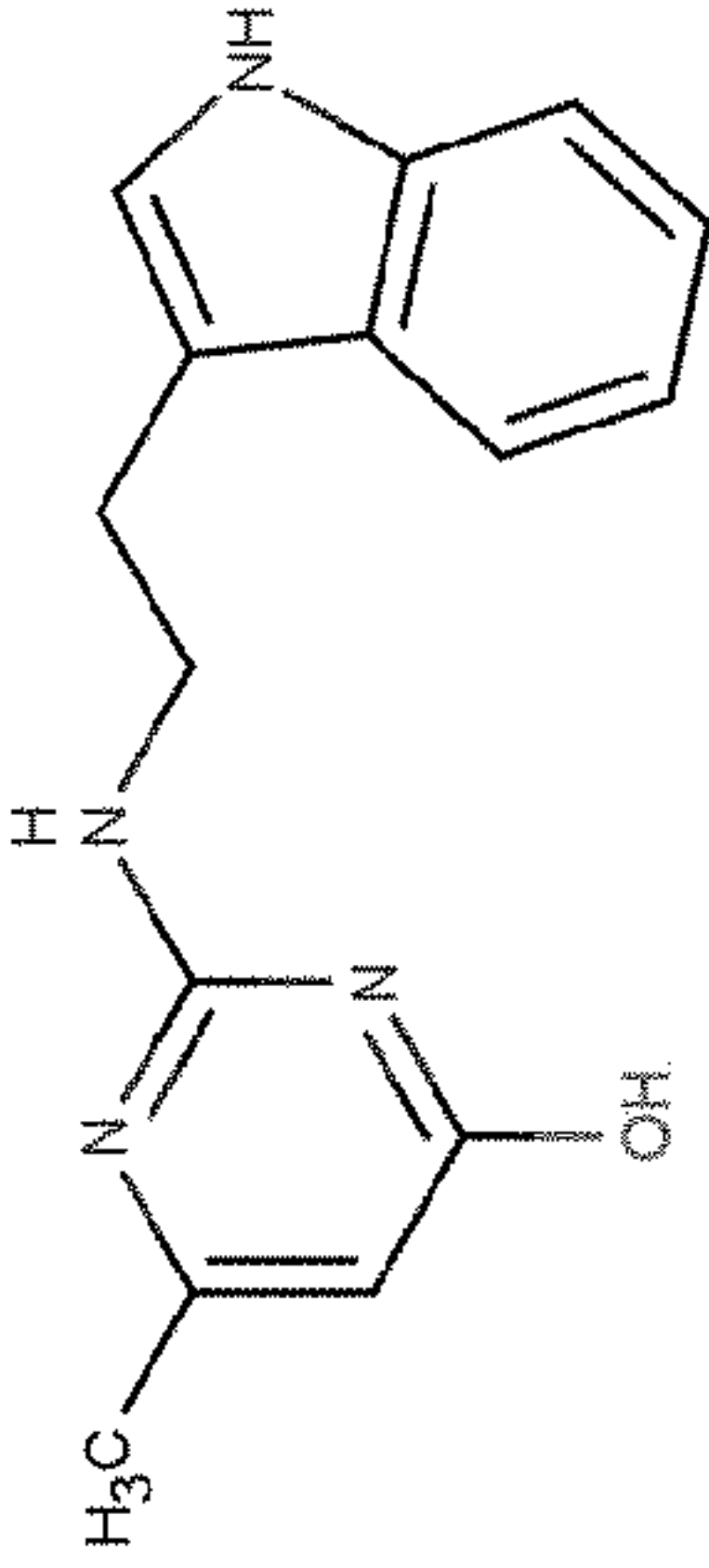
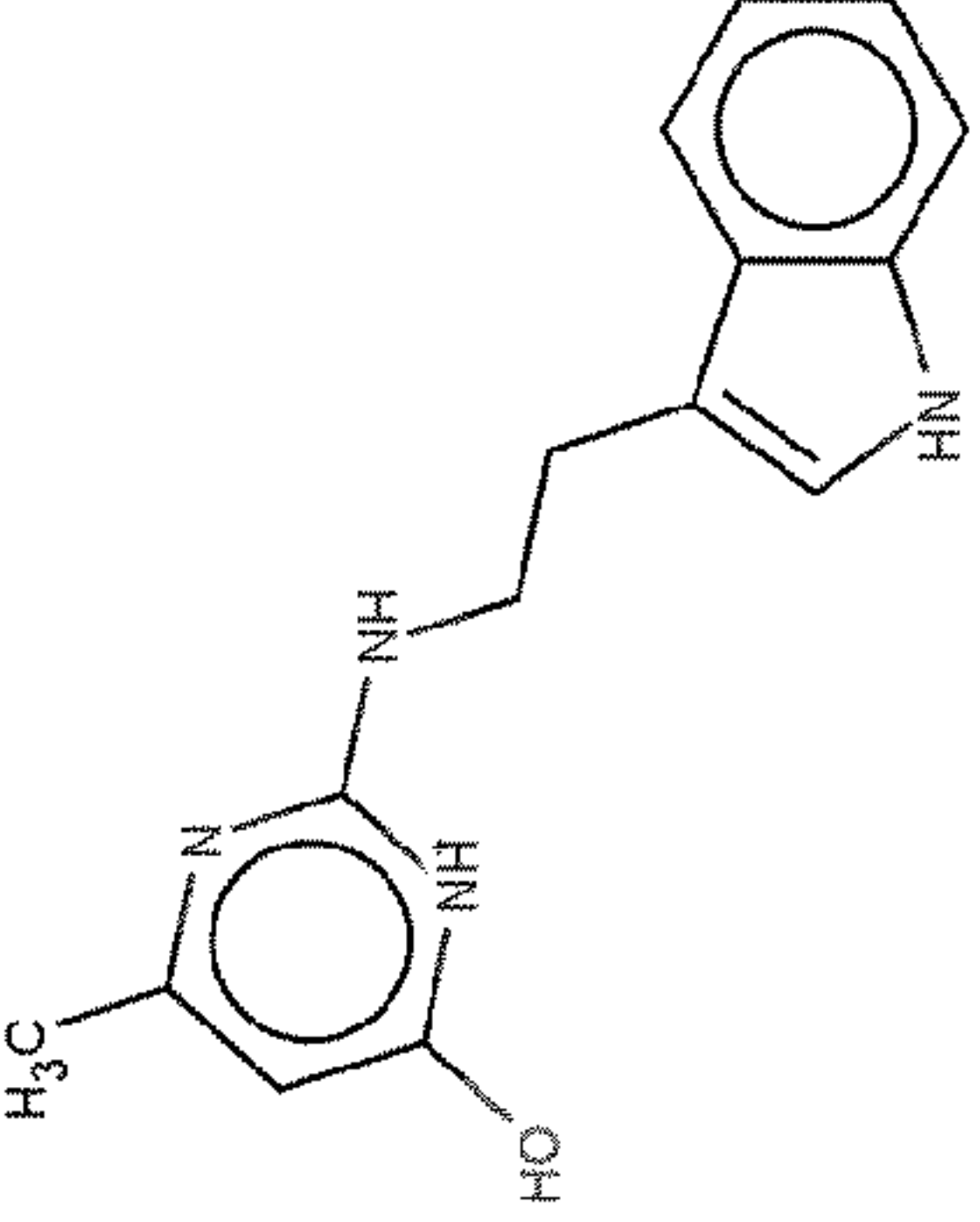
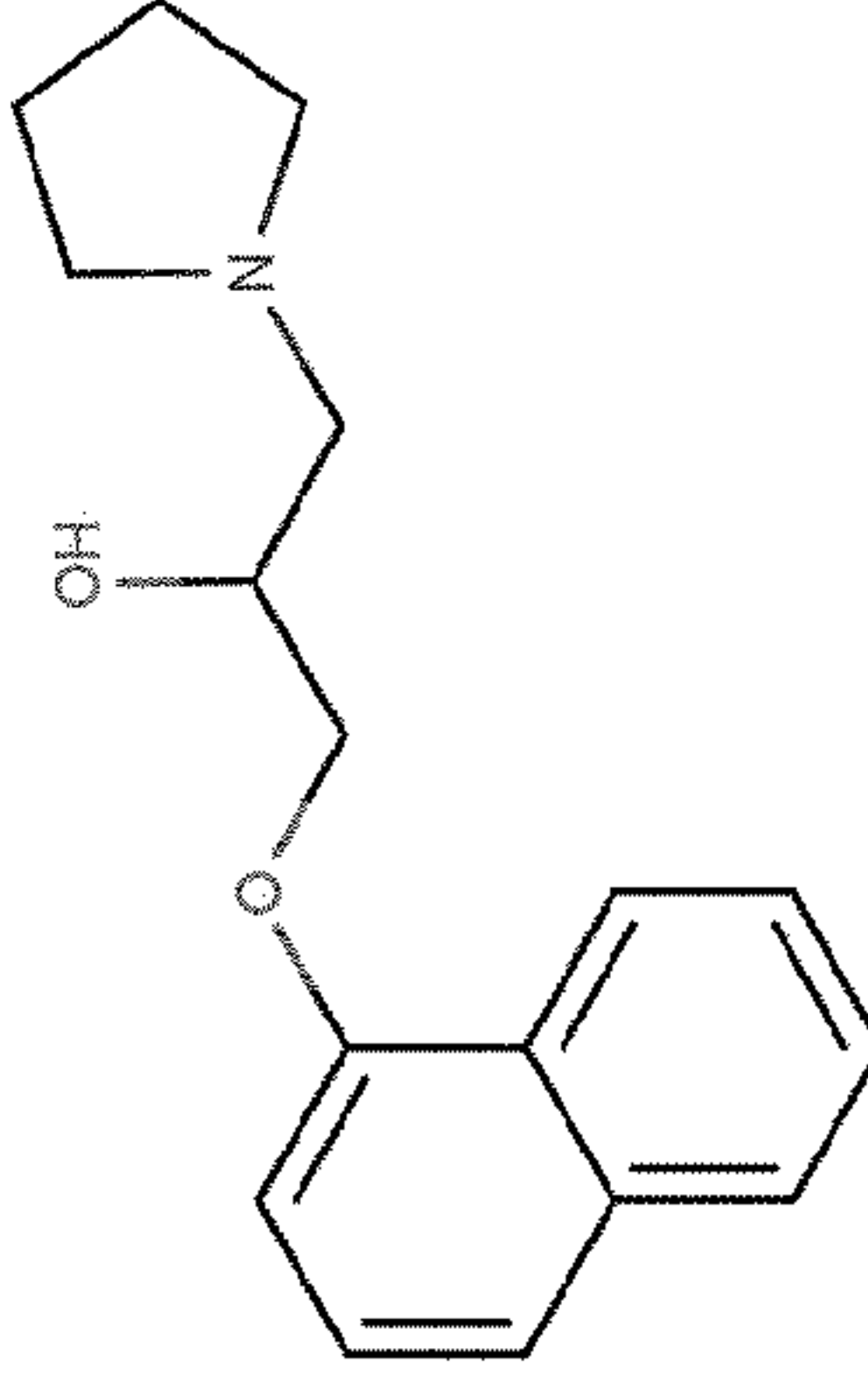
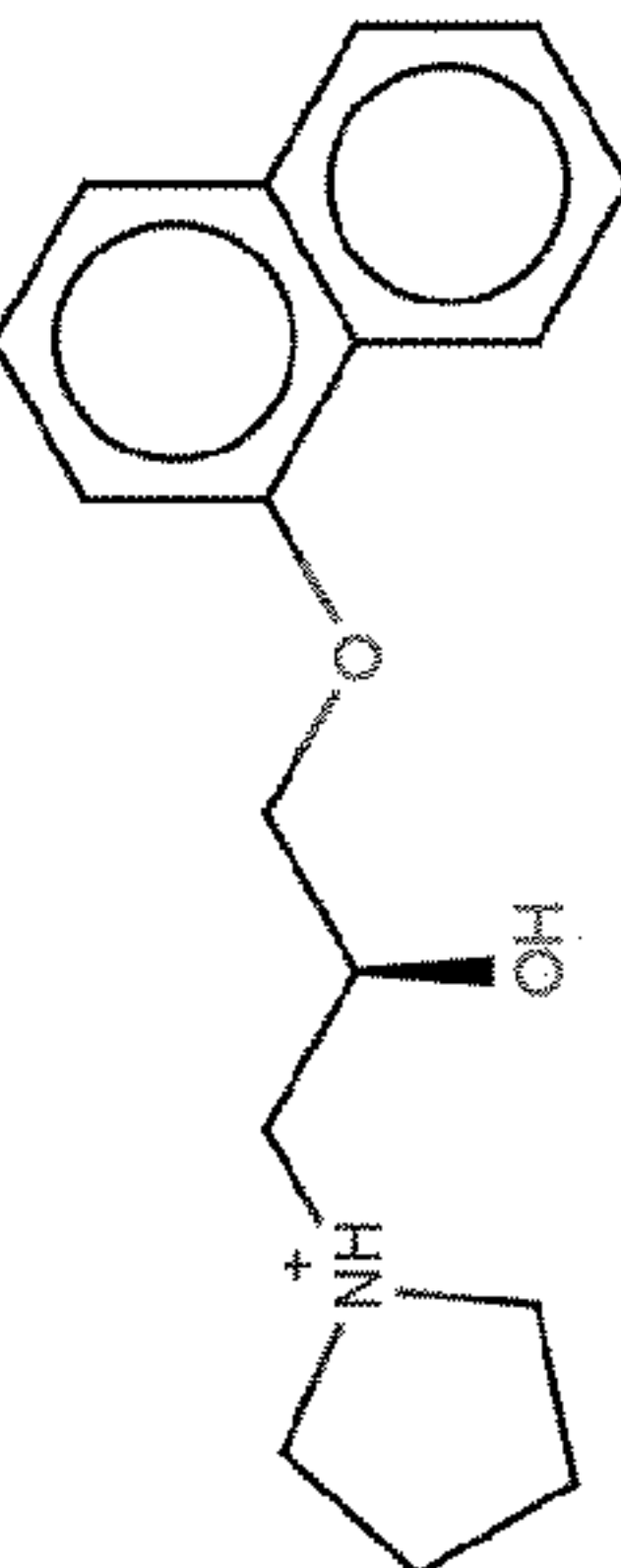
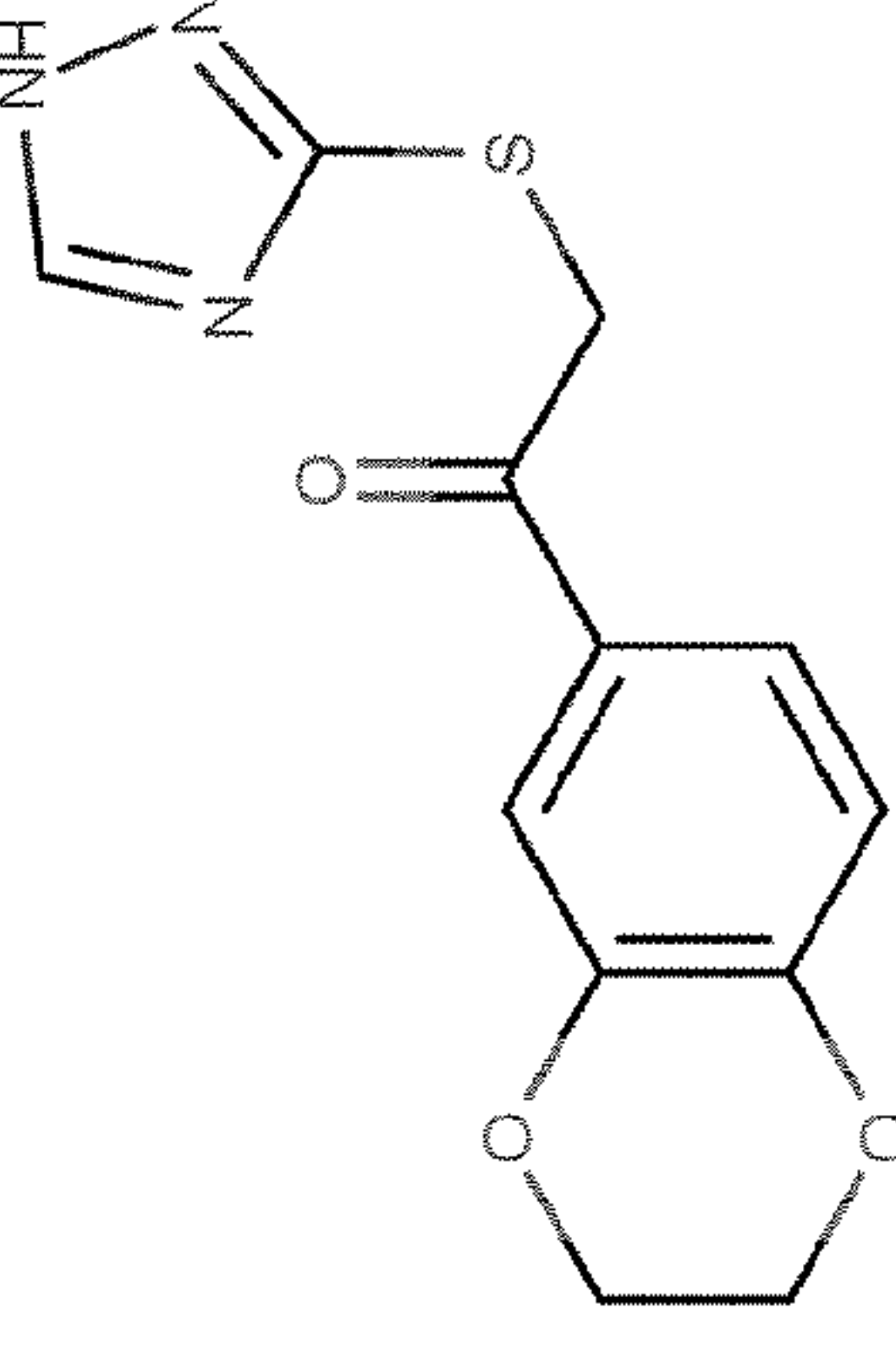
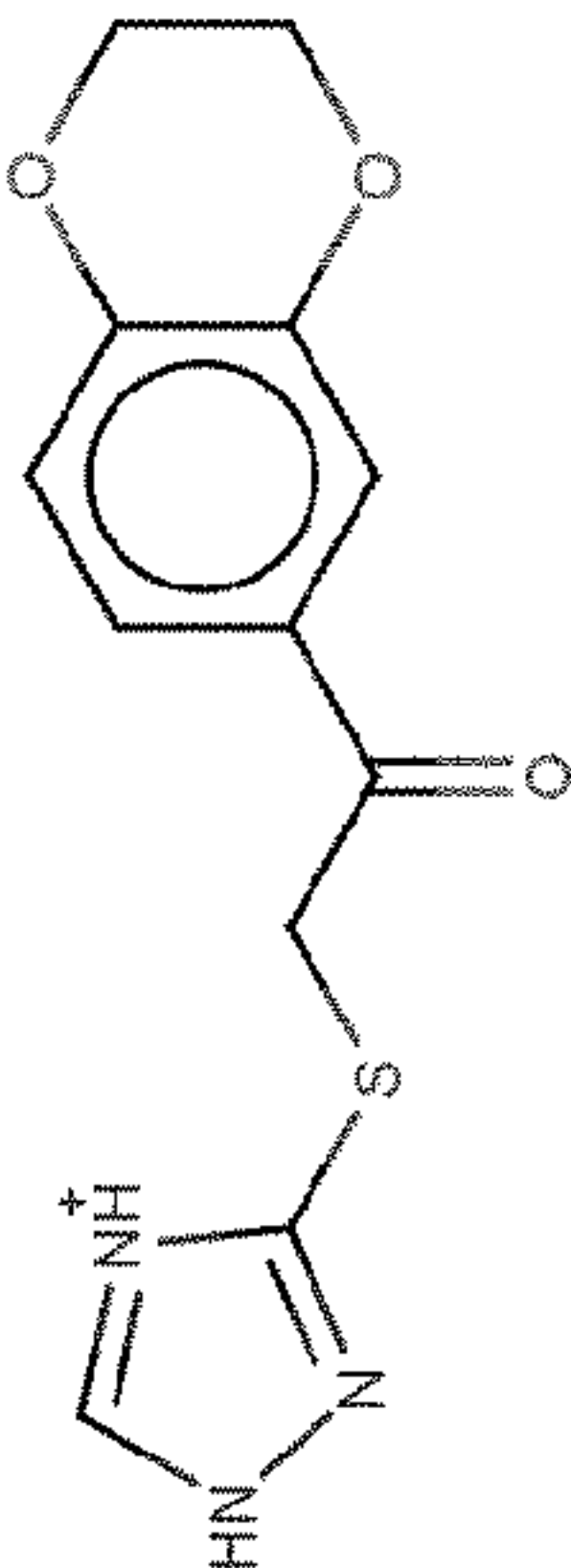
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 196		F3308-3248		F3308-3248_6_1 (1 of 12)
Compound 197		F3308-3604		F3308-3604_3_1 (1 of 14)
Compound 198		F3314-0051		F3314-0051_1 (1 of 16)

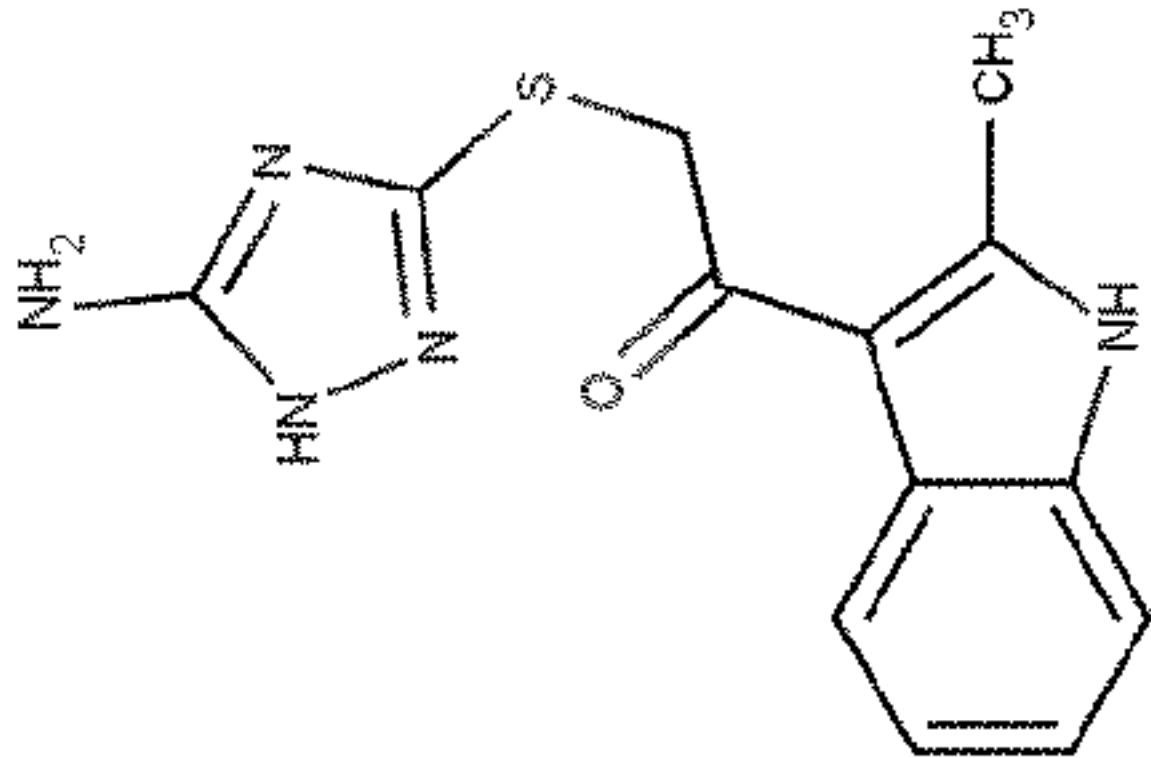
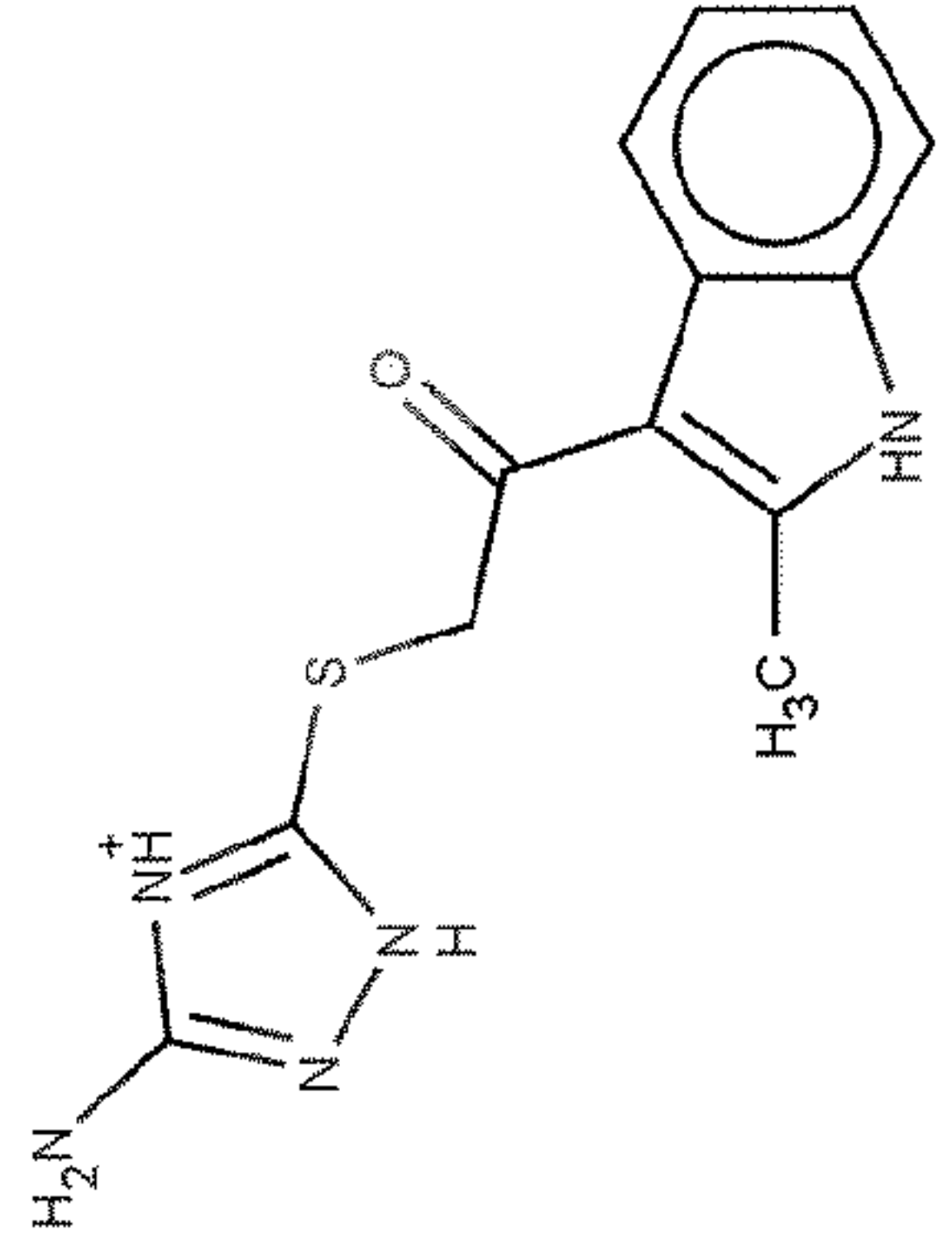
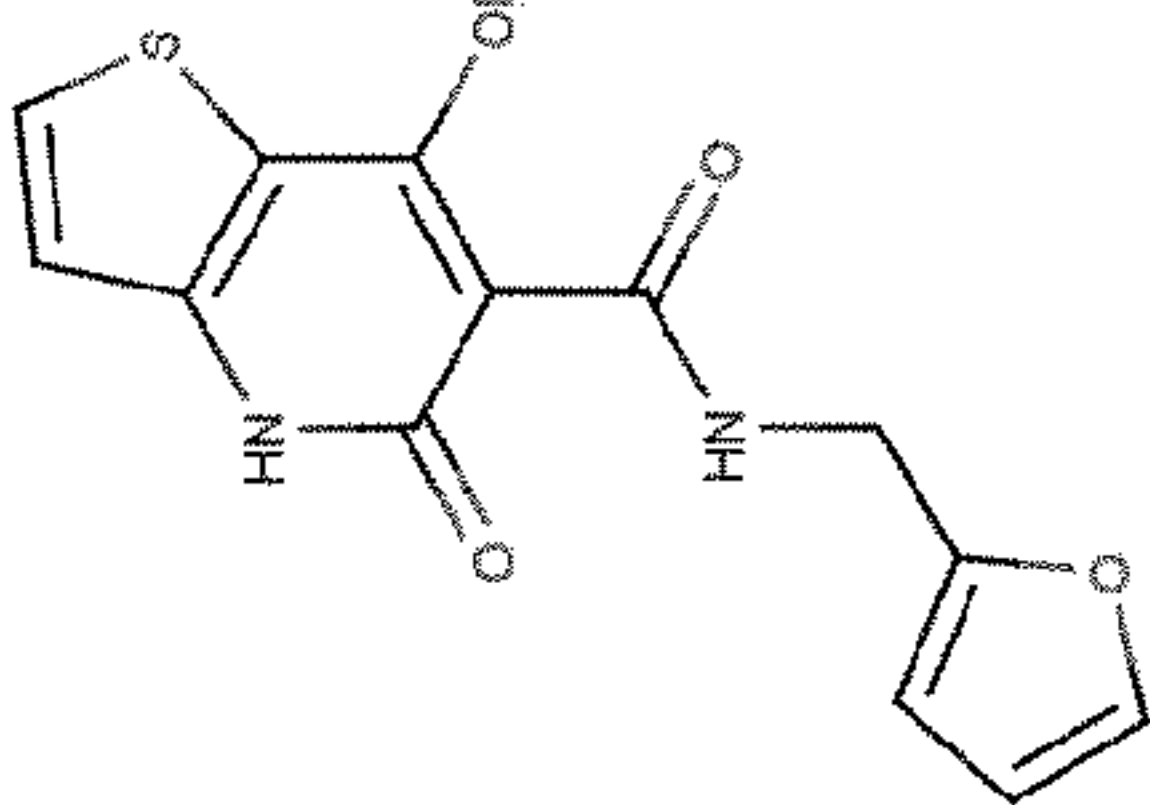
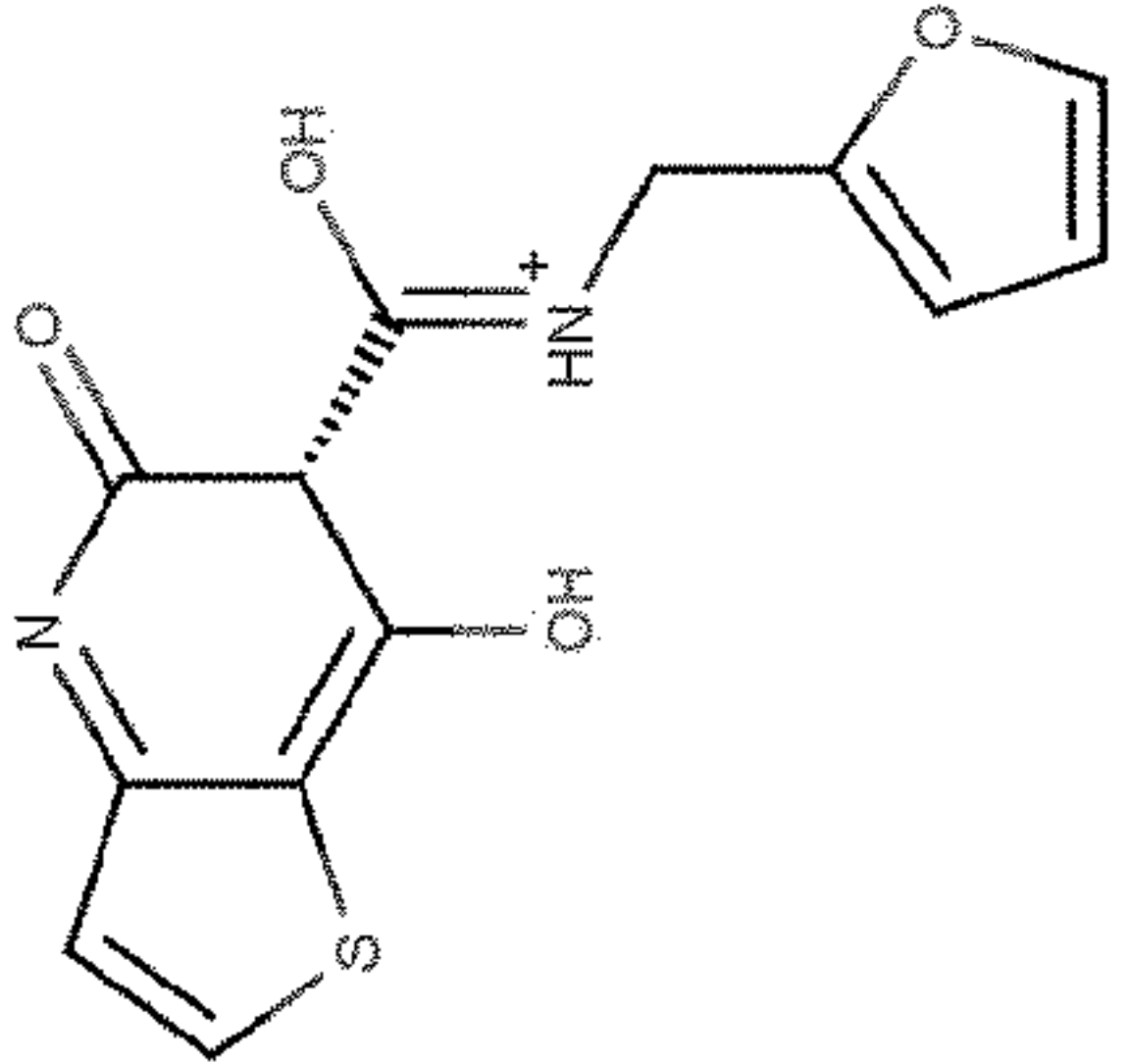
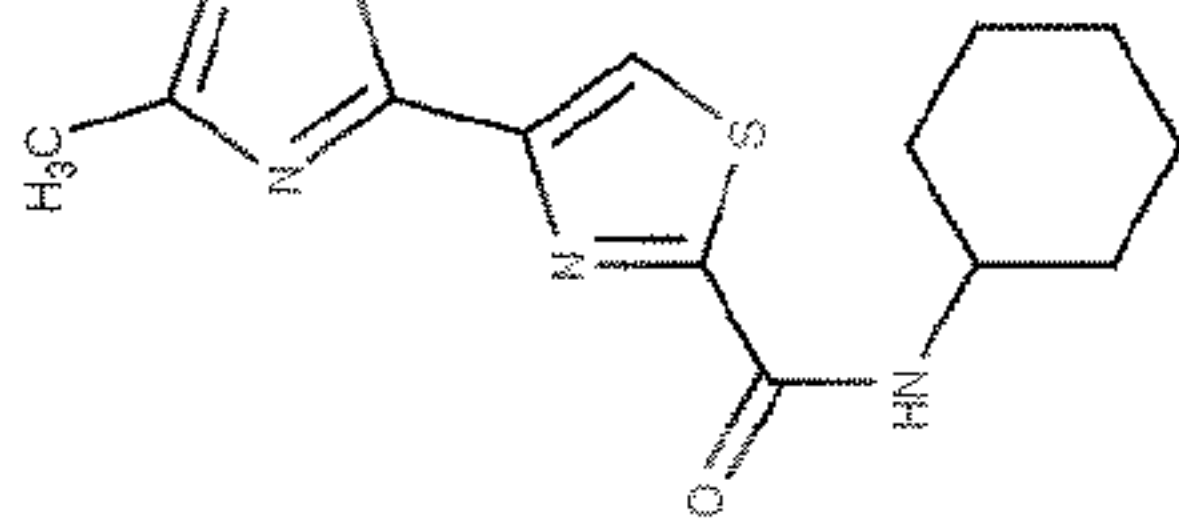
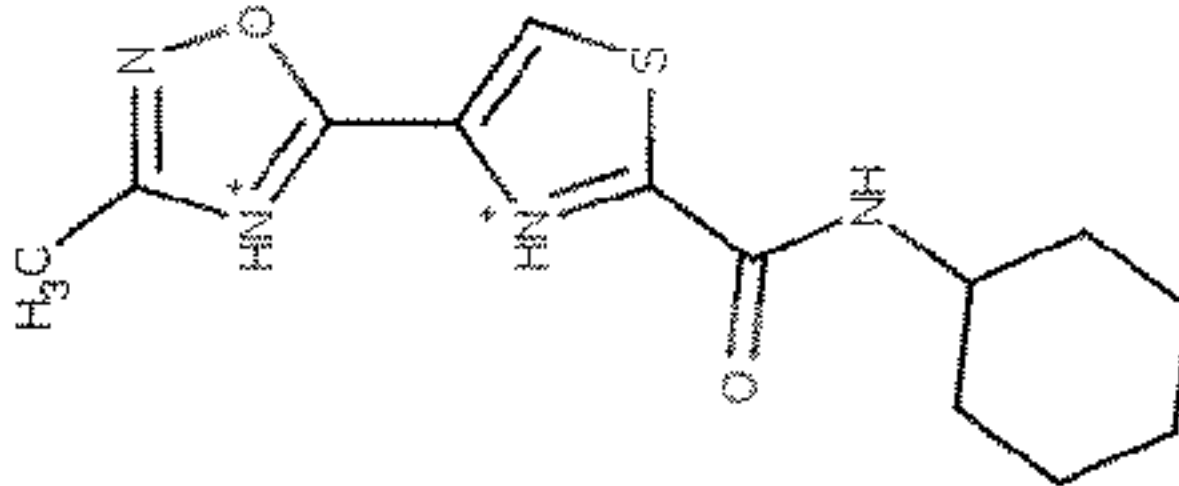
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 199		F3314-0053		F3314-0053_1_3 (1 of 8)
Compound 200		F3314-0059		F3314-0059_2 (1 of 4)
Compound 201		F3314-0071		F3314-0071_2 (1 of 4)

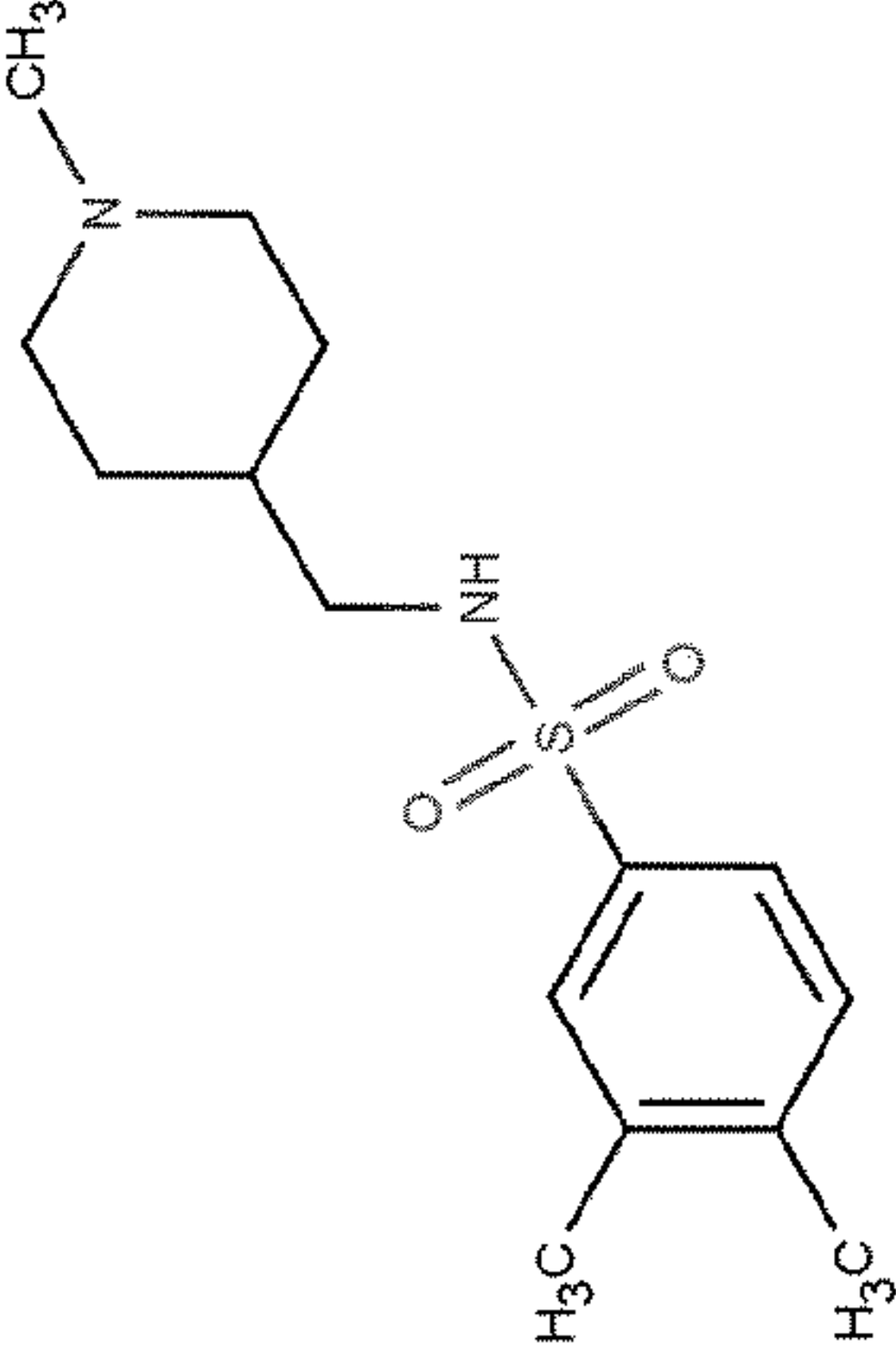
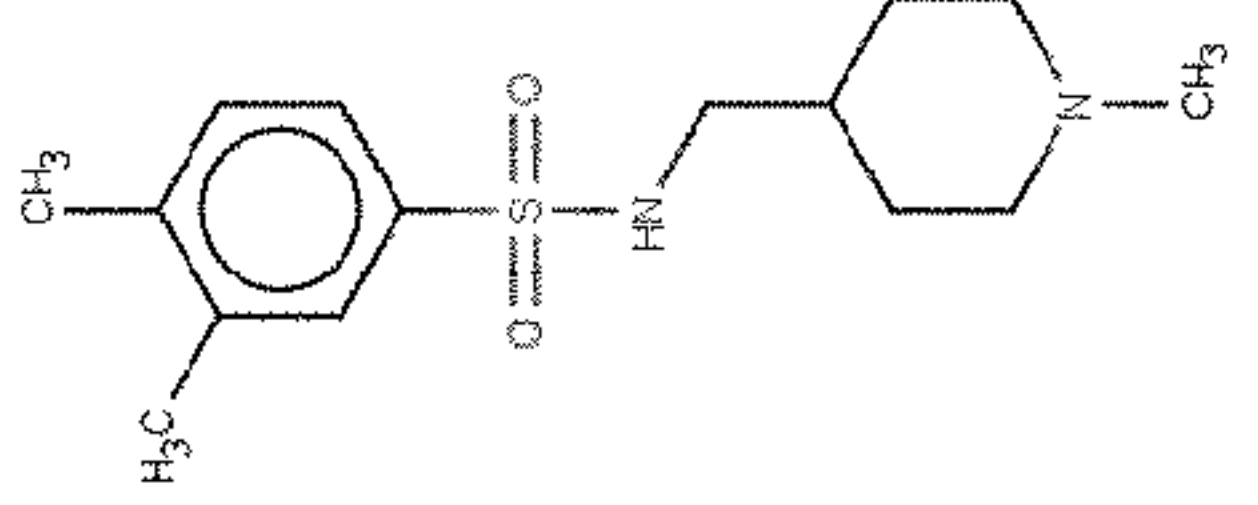
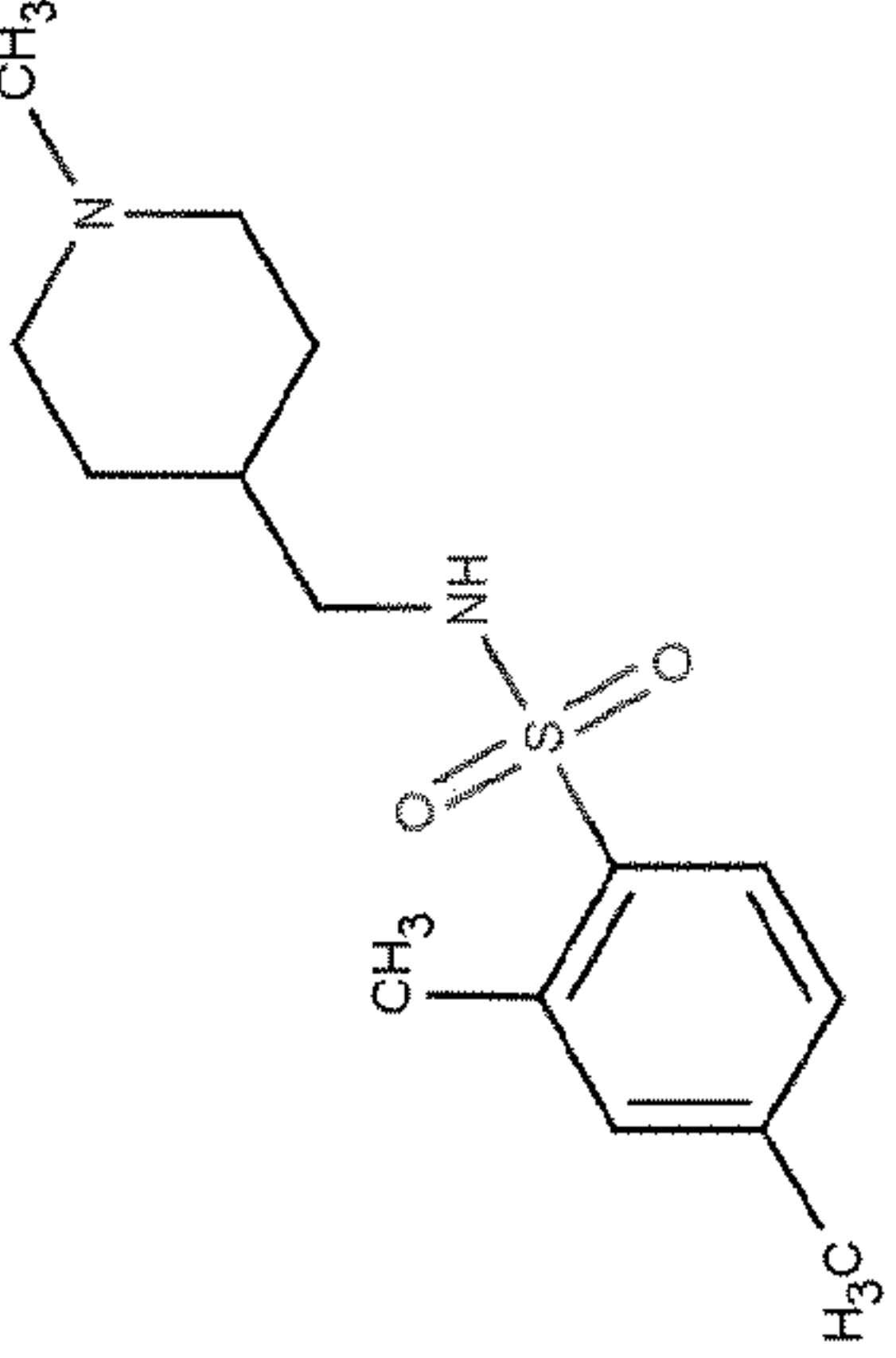
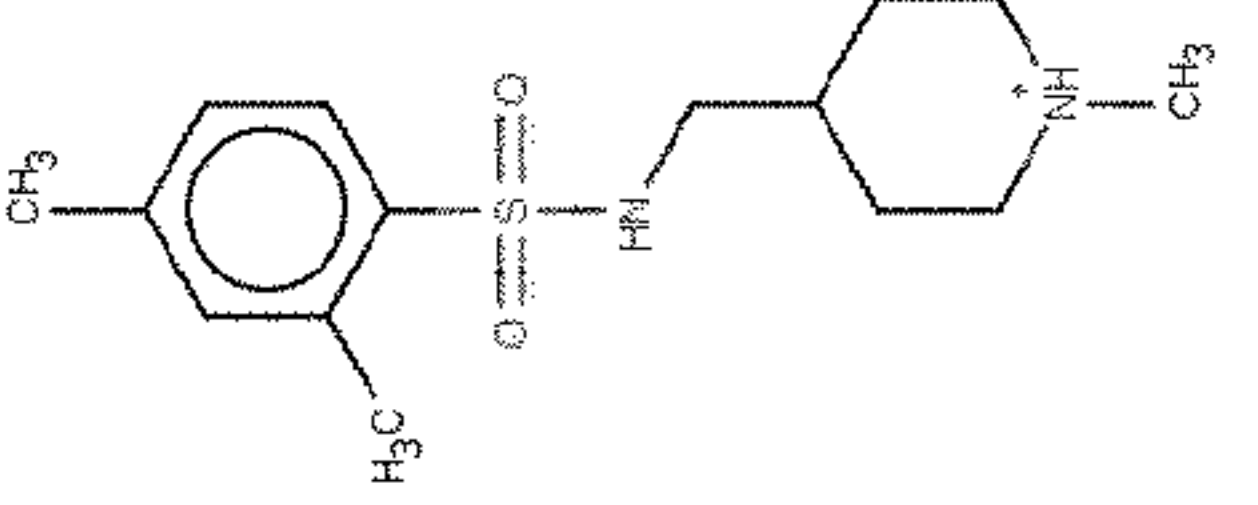
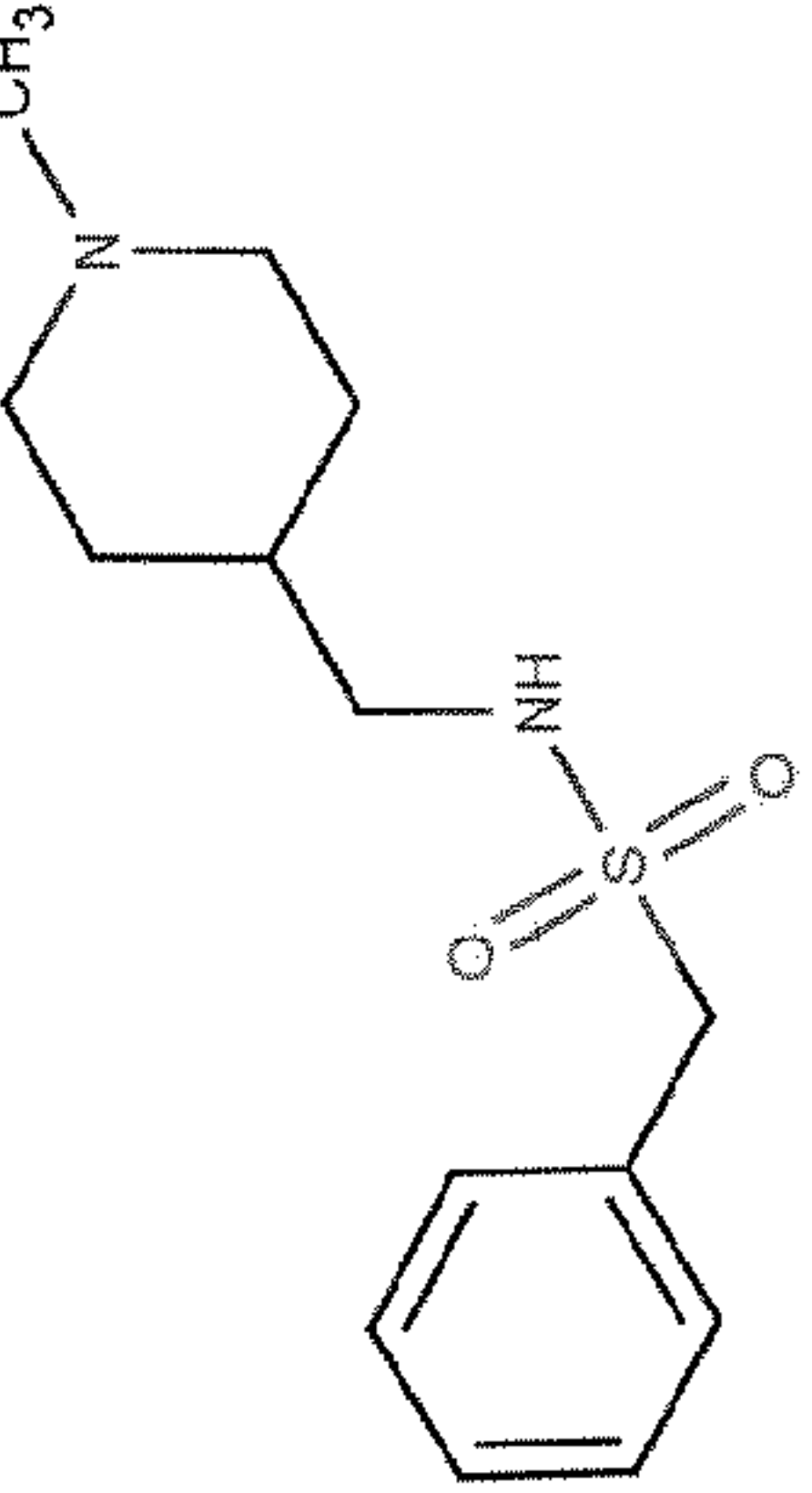
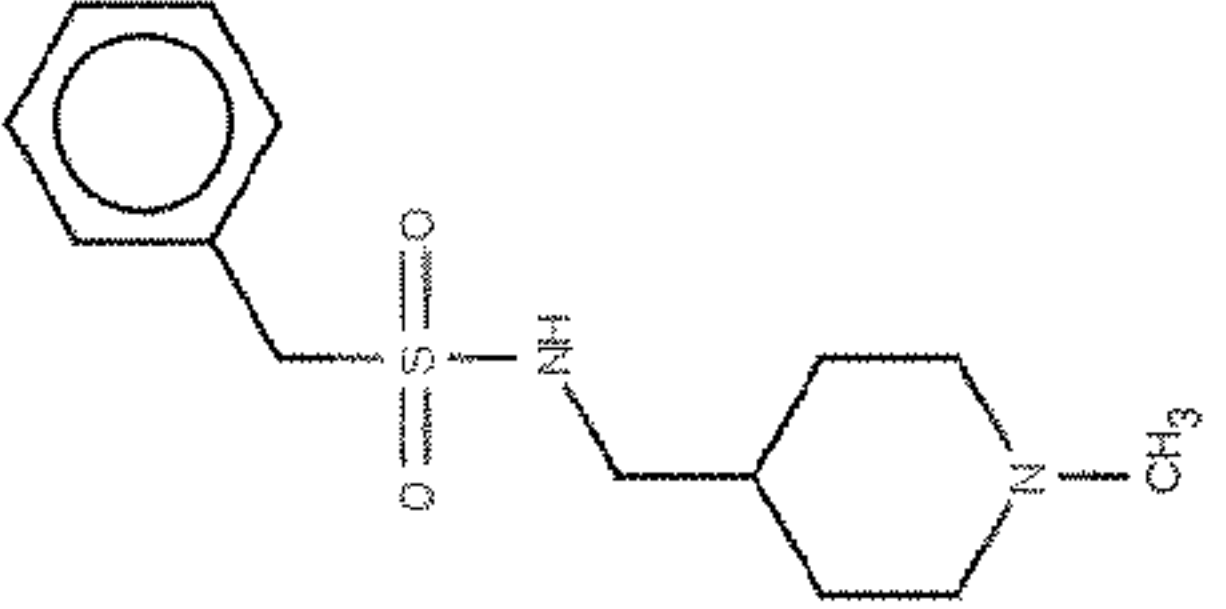
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 202		F3316-0057		F3316-0057_1_2 (1 of 4)
Compound 203		F3316-0139		F3316-0139_3_2 (1 of 8)
Compound 204		F3316-0145		F3316-0145_1_2 (1 of 4)

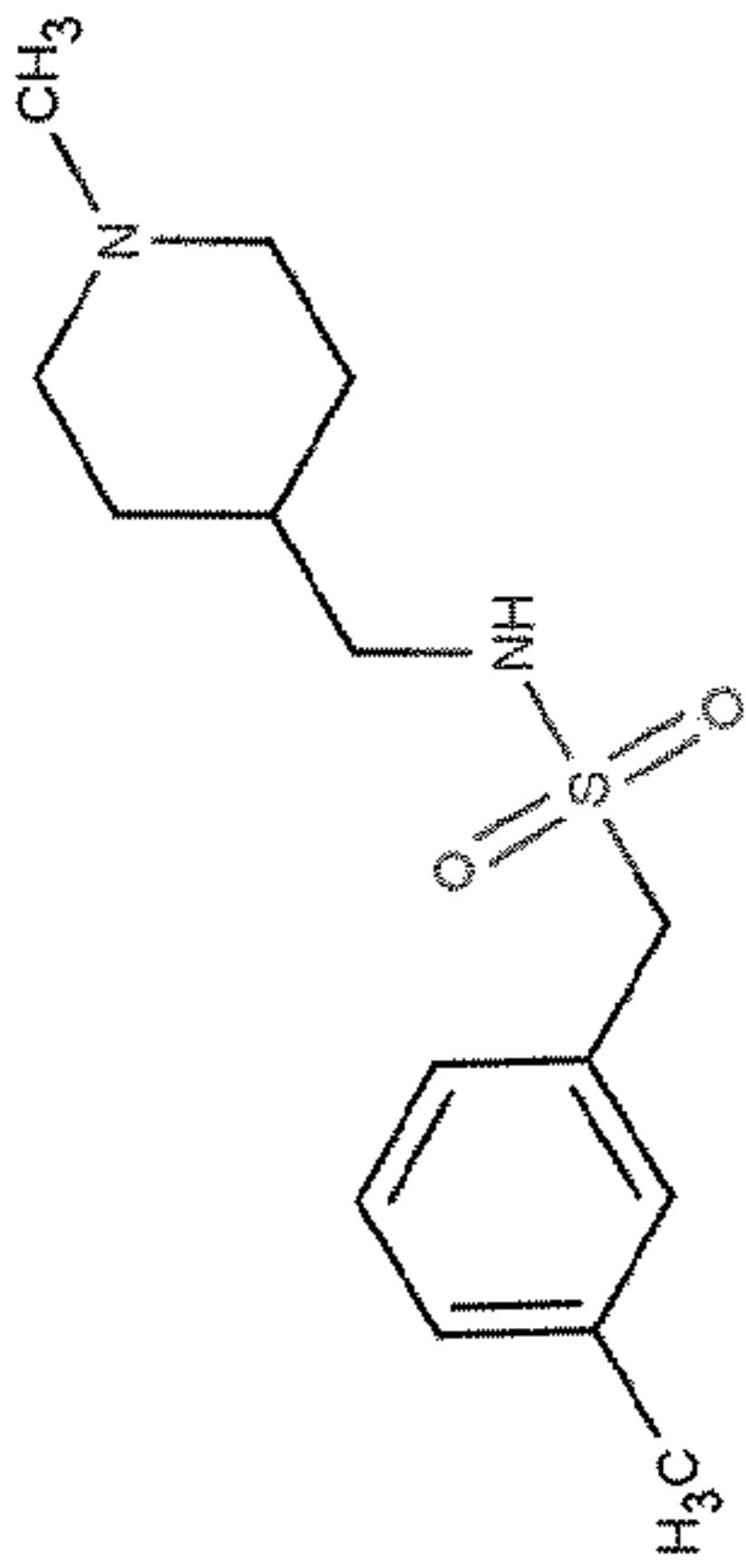
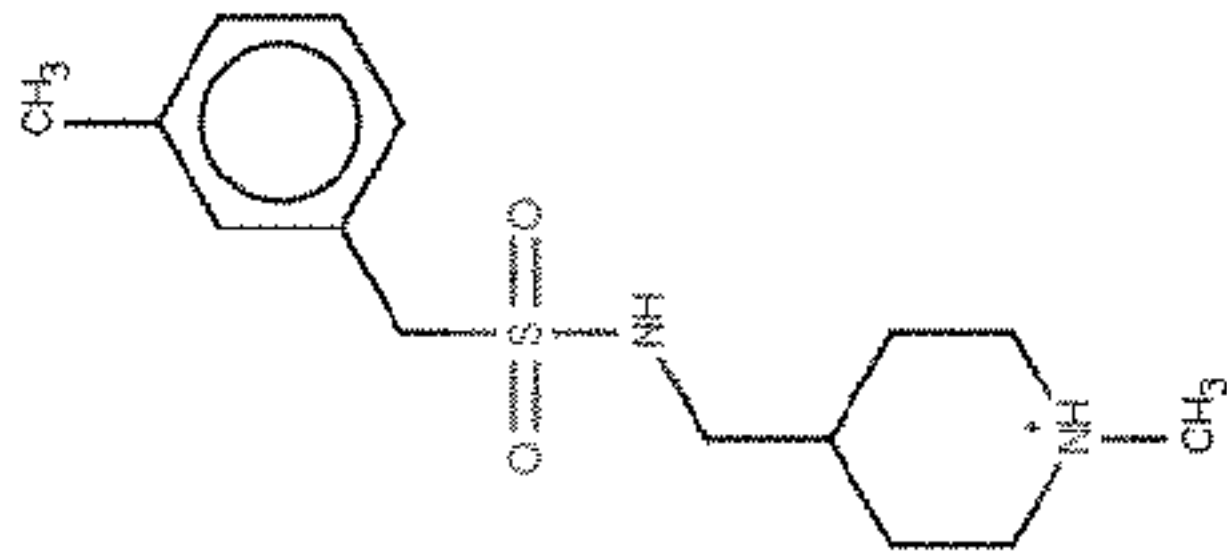
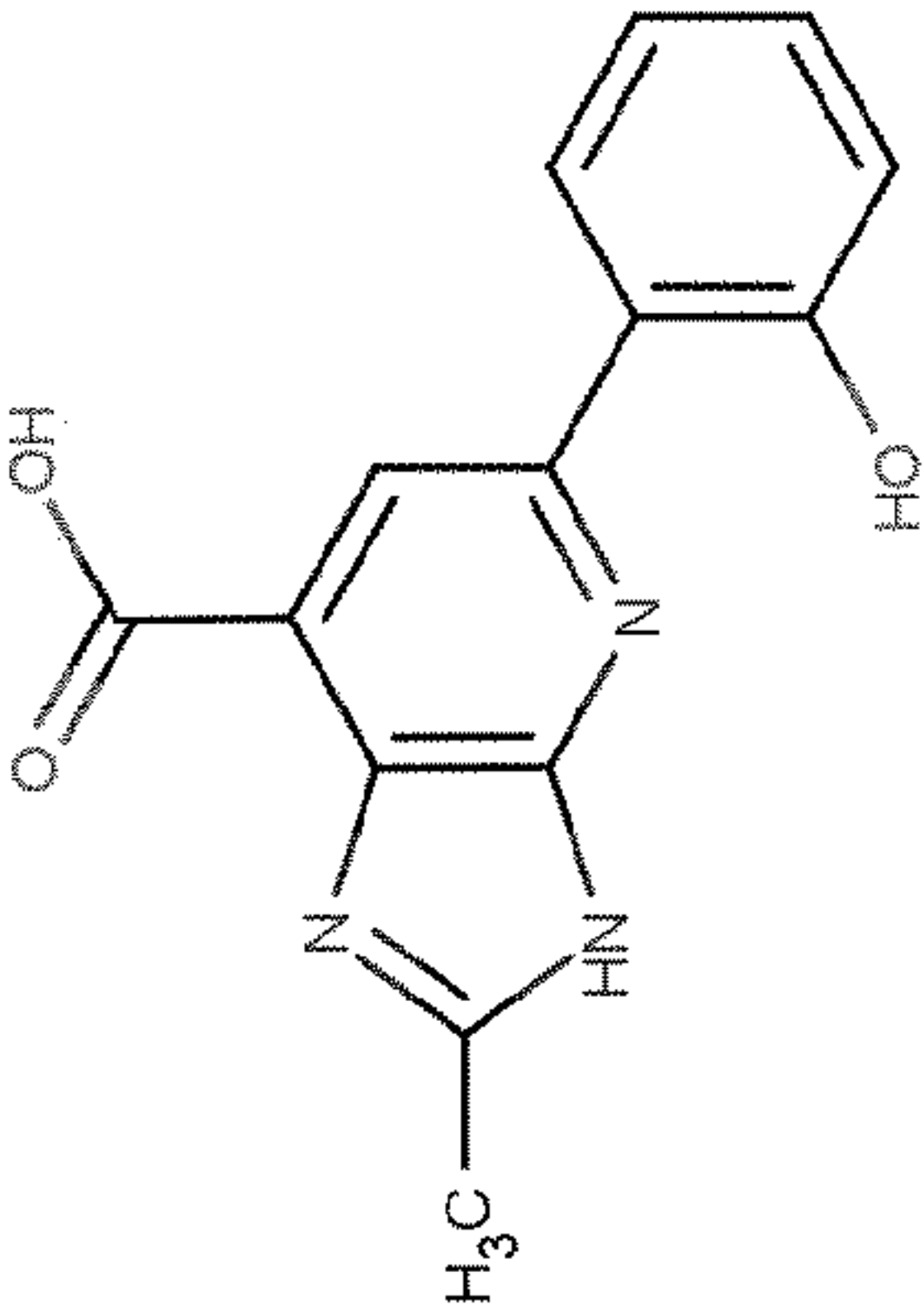
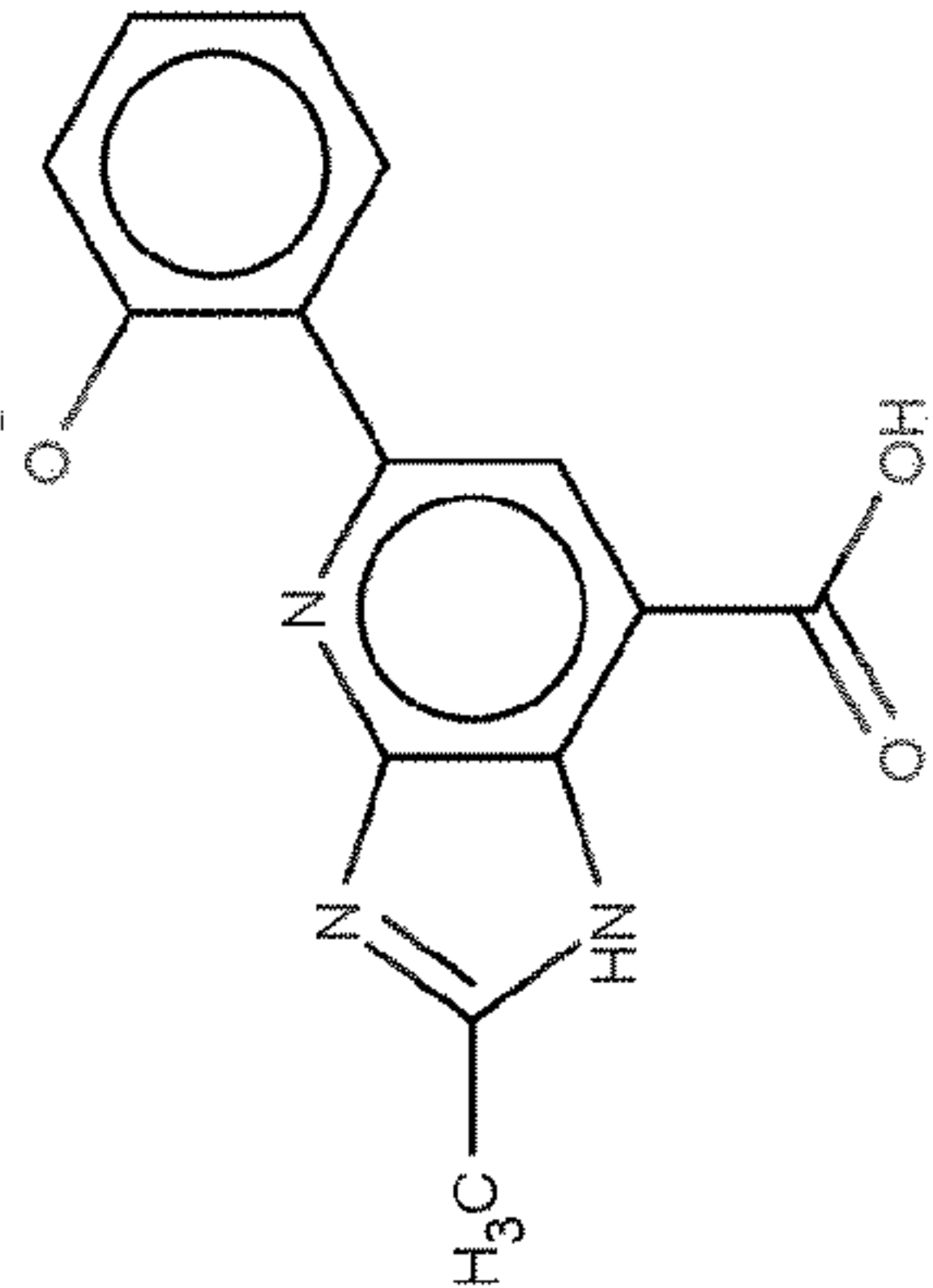
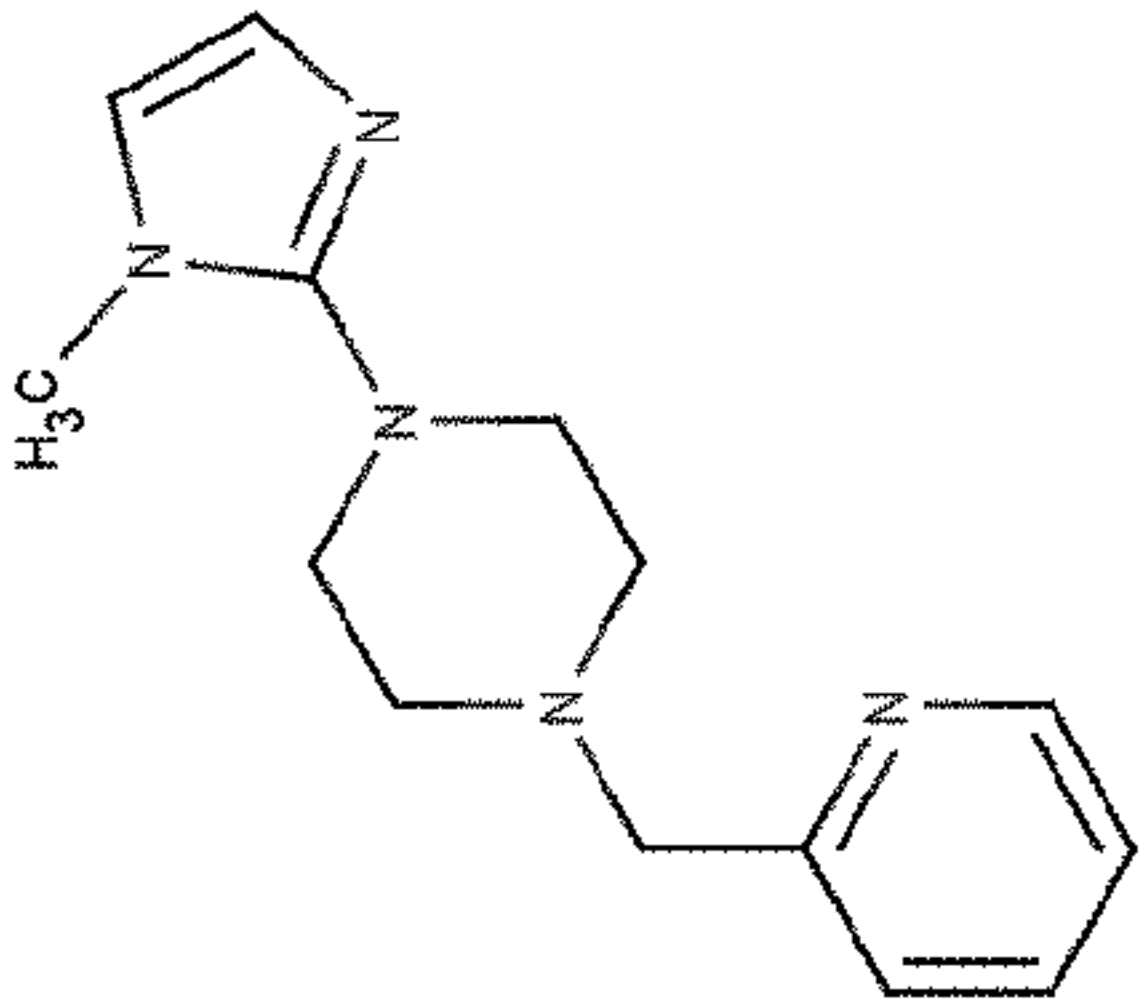
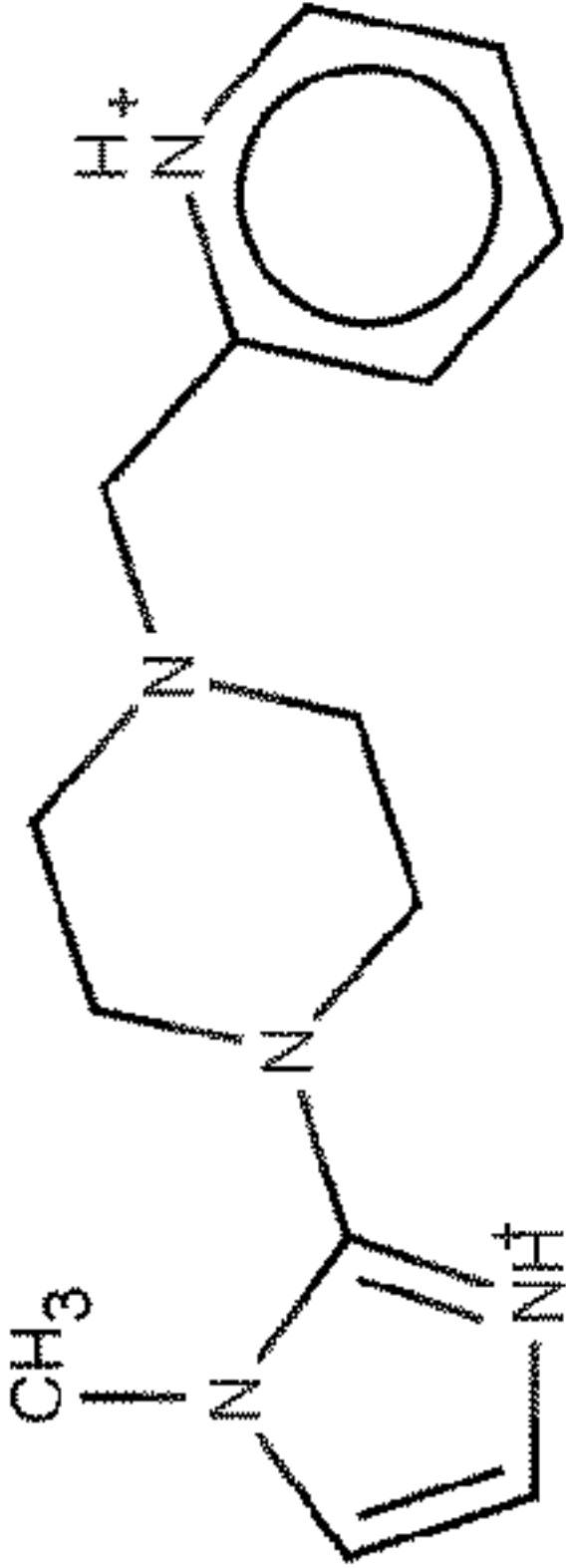
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 205		F3316-0146		F3316-0146_1_2 (1 of 4)
Compound 206		F3316-0166		F3316-0166_1_1 (1 of 4)
Compound 207		F3320-0301		F3320-0301_5_1 (1 of 7)

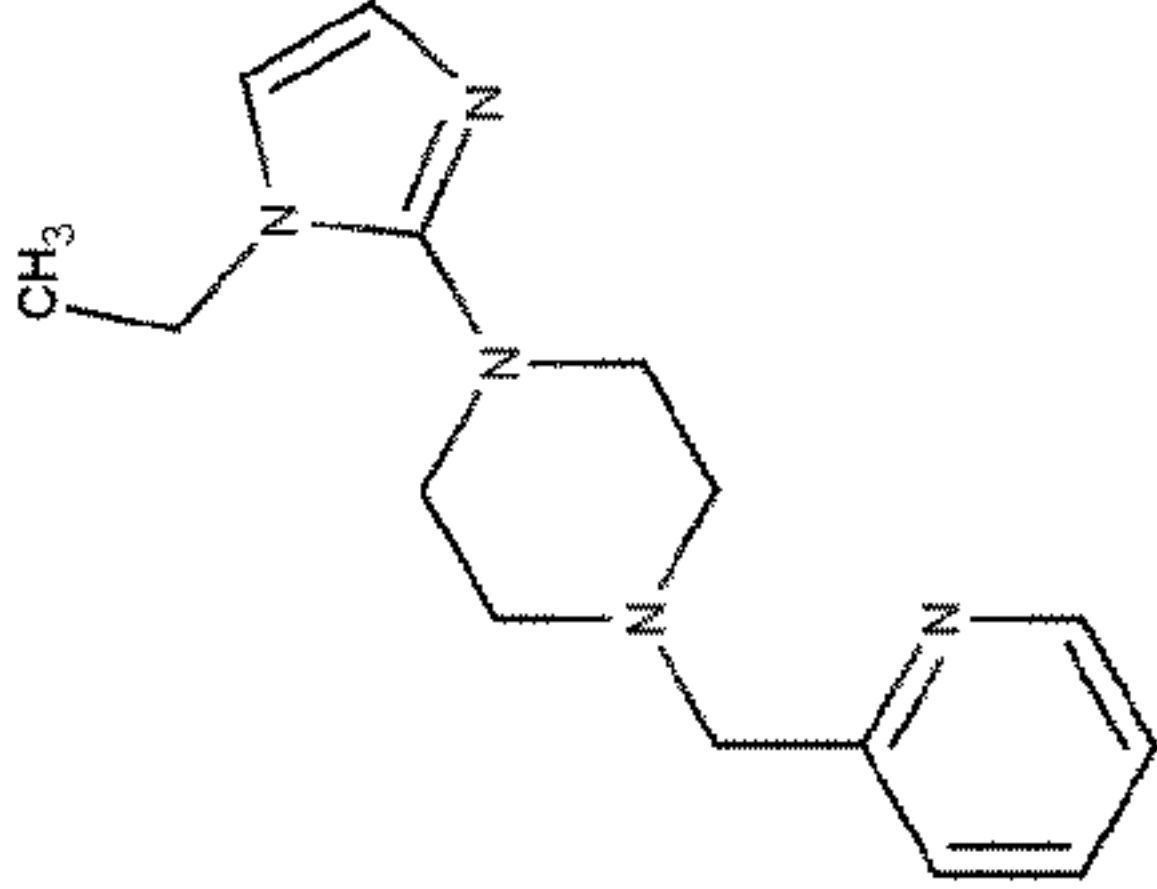
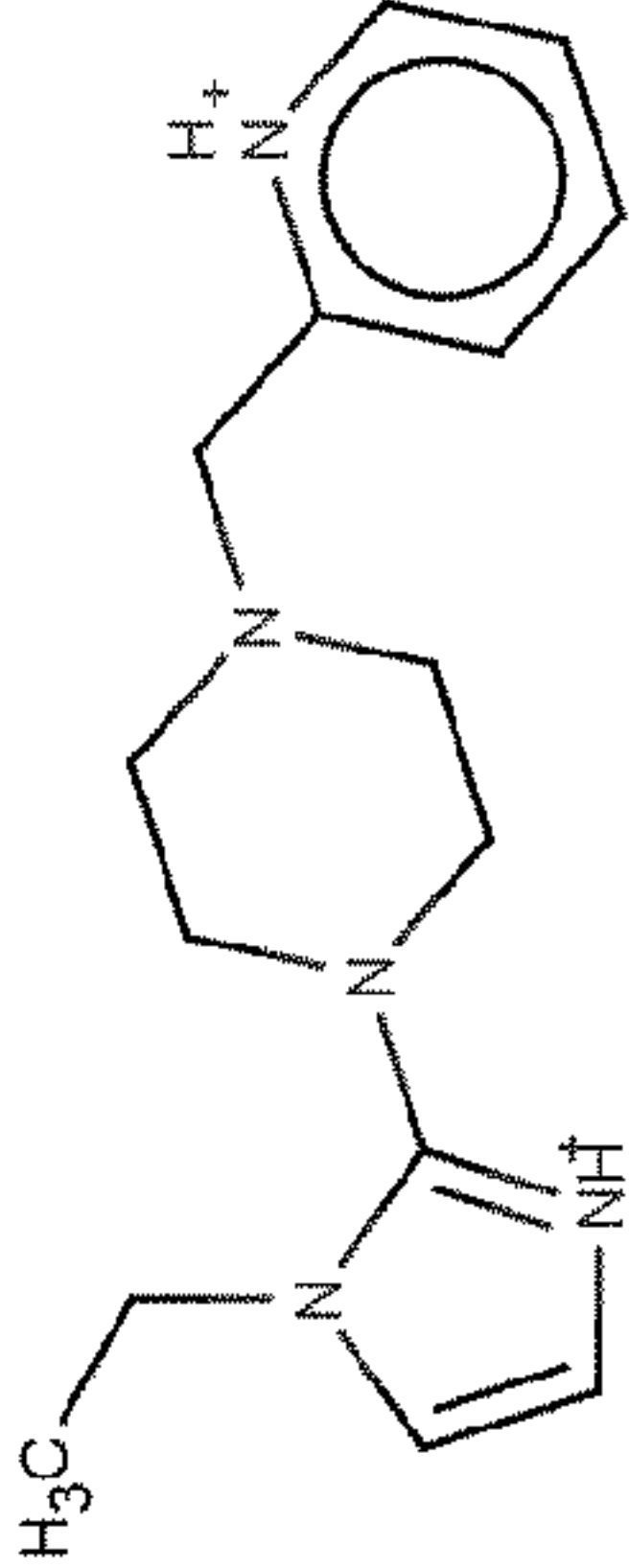
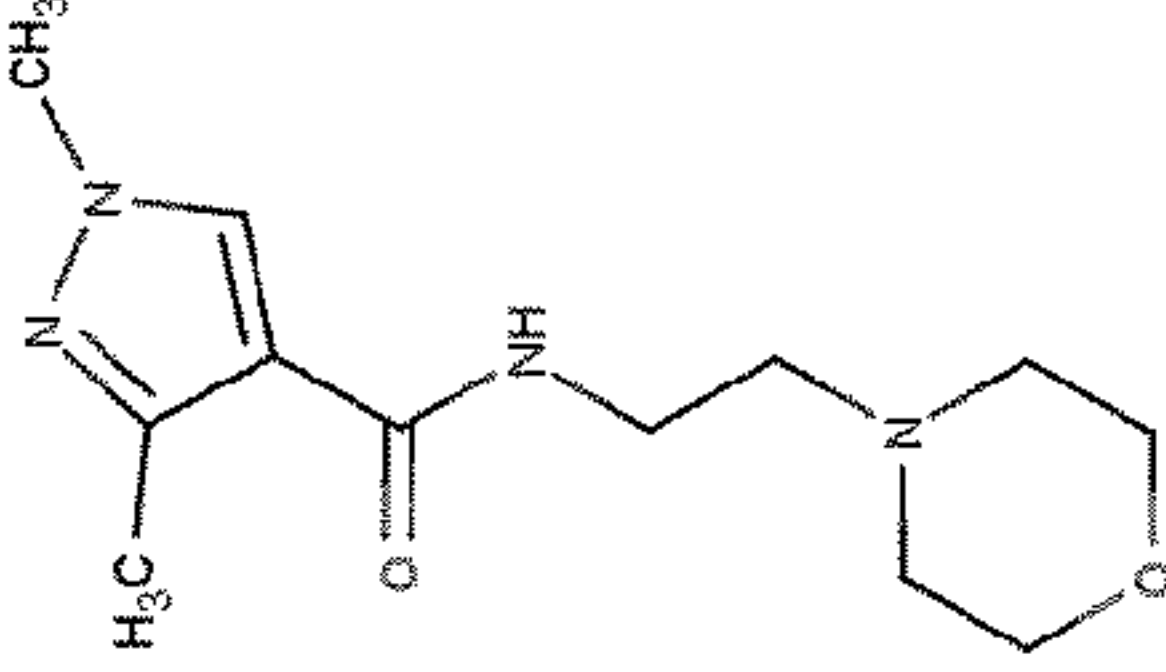
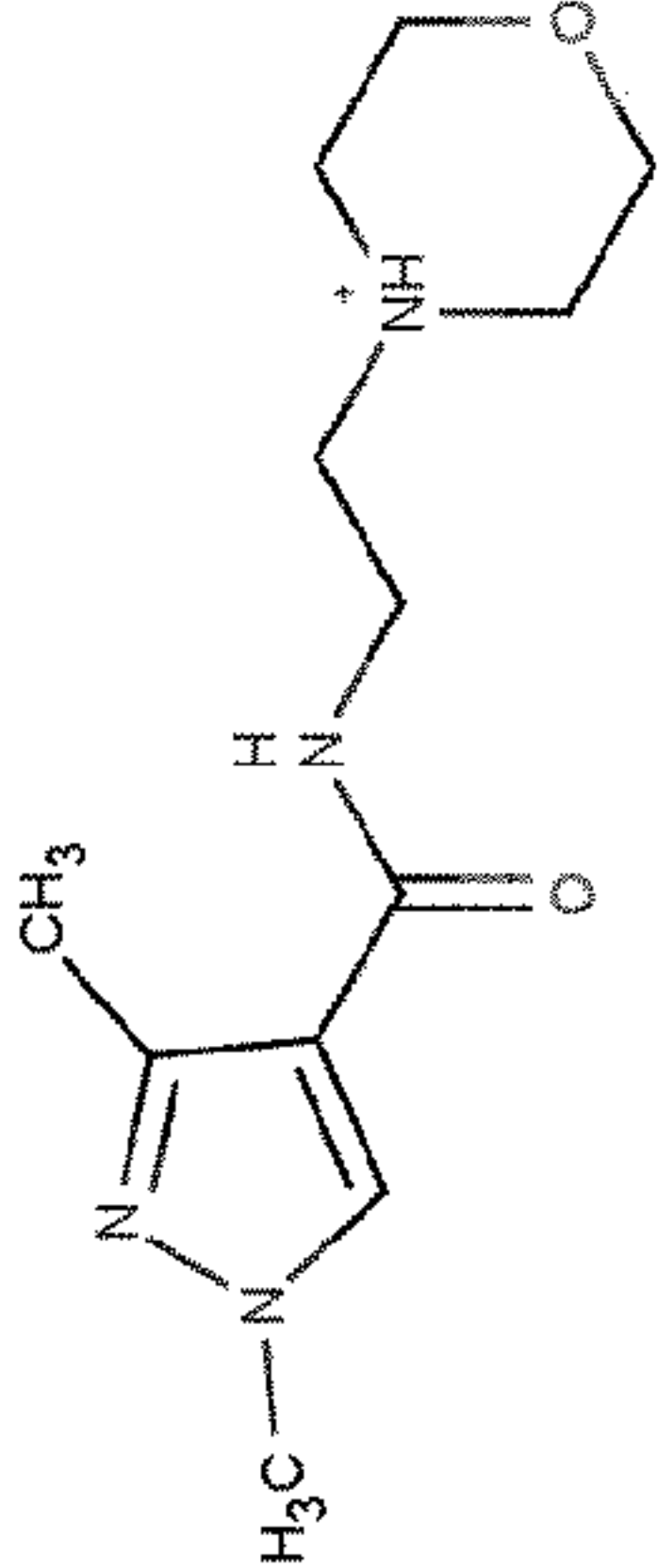
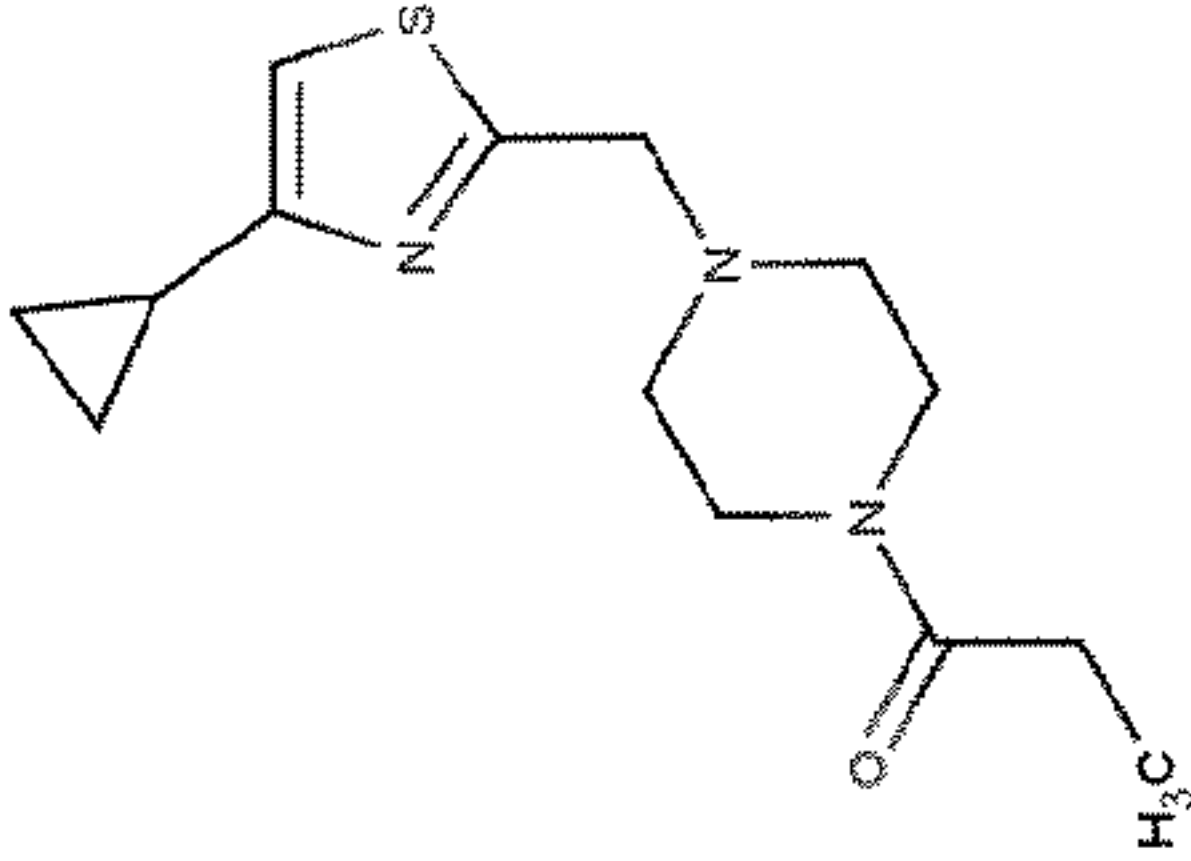
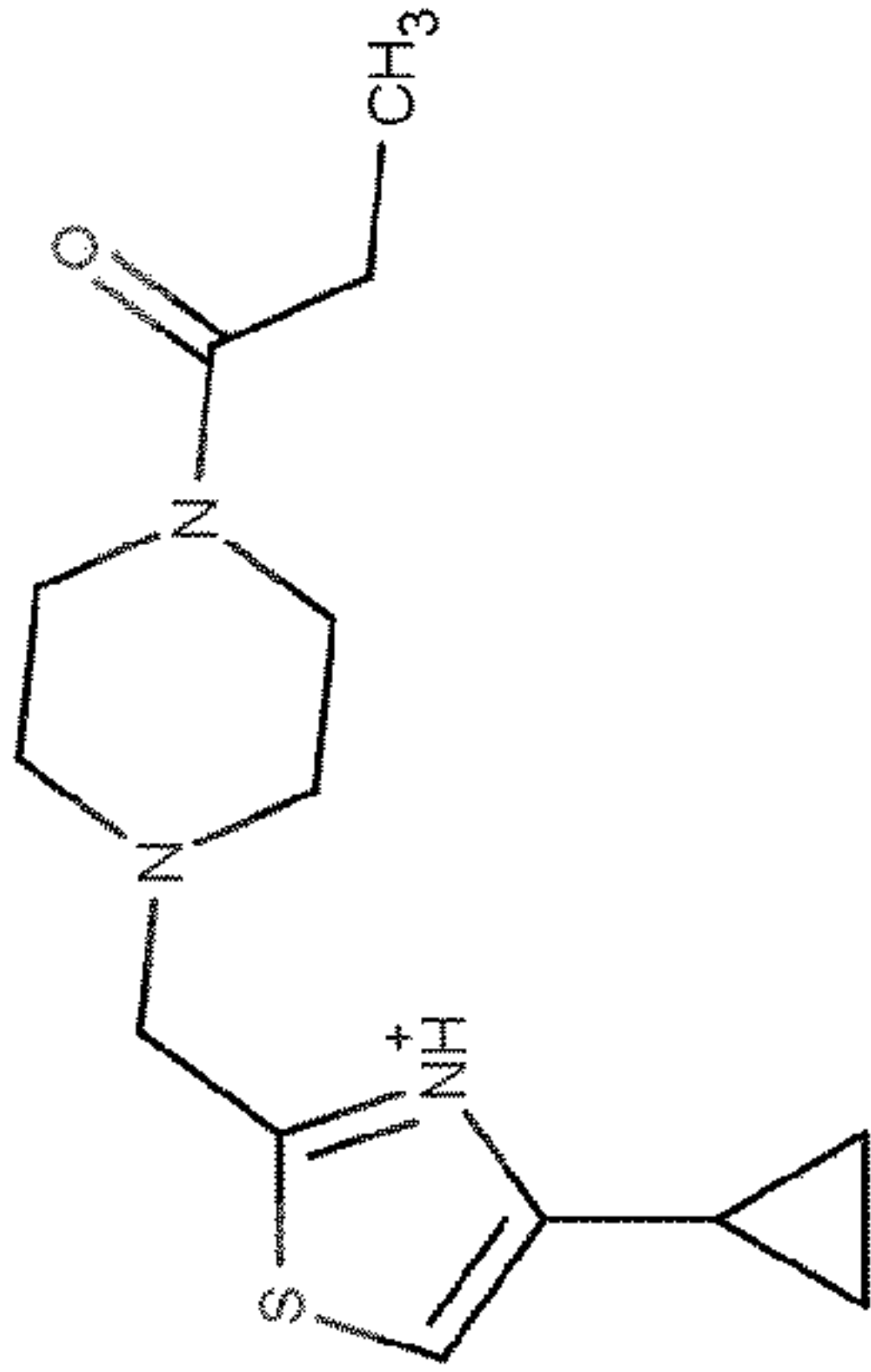
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 208		F3331-0647		F3331-0647_1_1 (1 of 4)
Compound 209		F3350-0739		F3350-0739_1_1 (1 of 4)
Compound 210		F3358-0351		F3358-0351_18_2 (1 of 26)

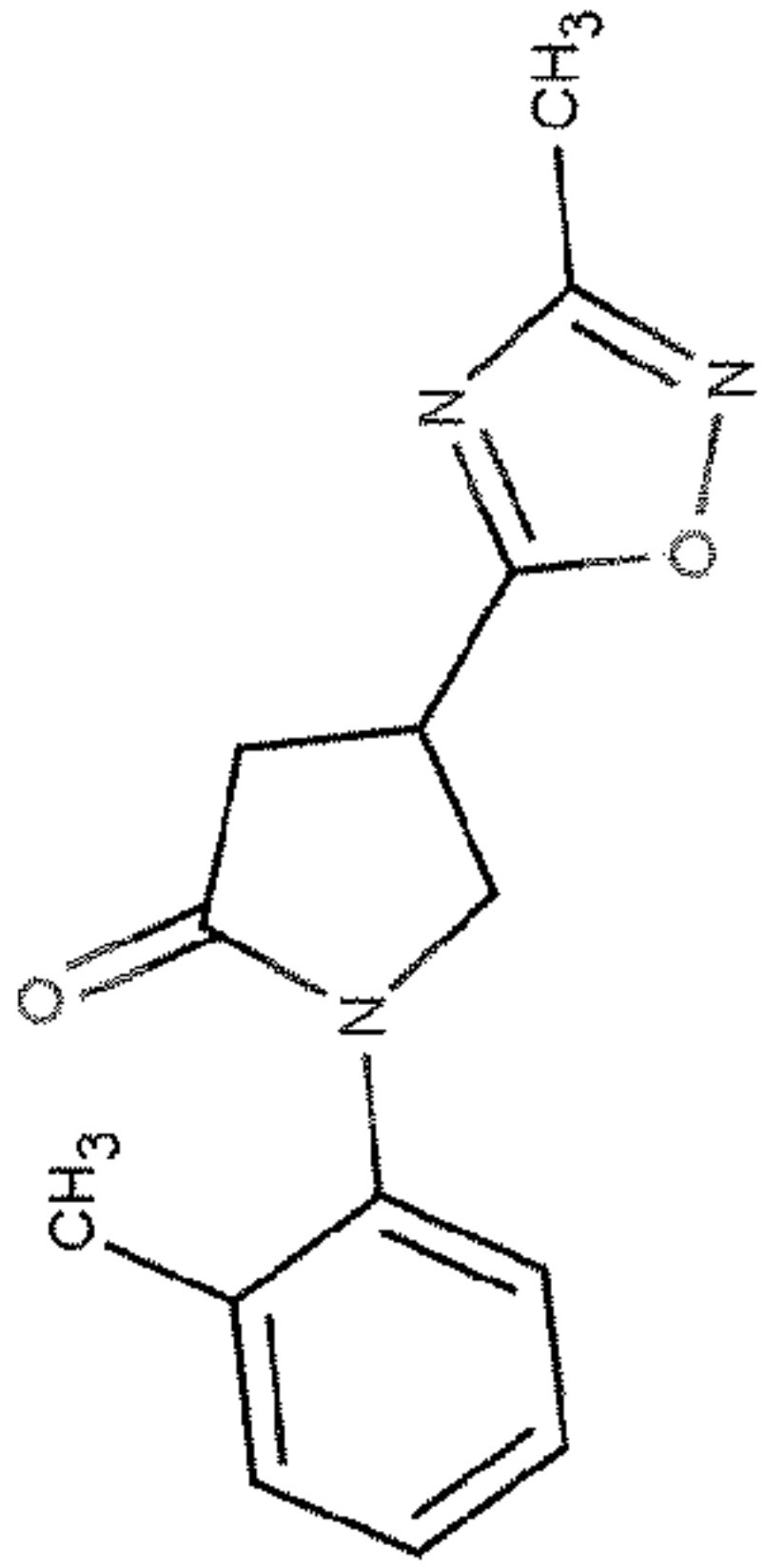
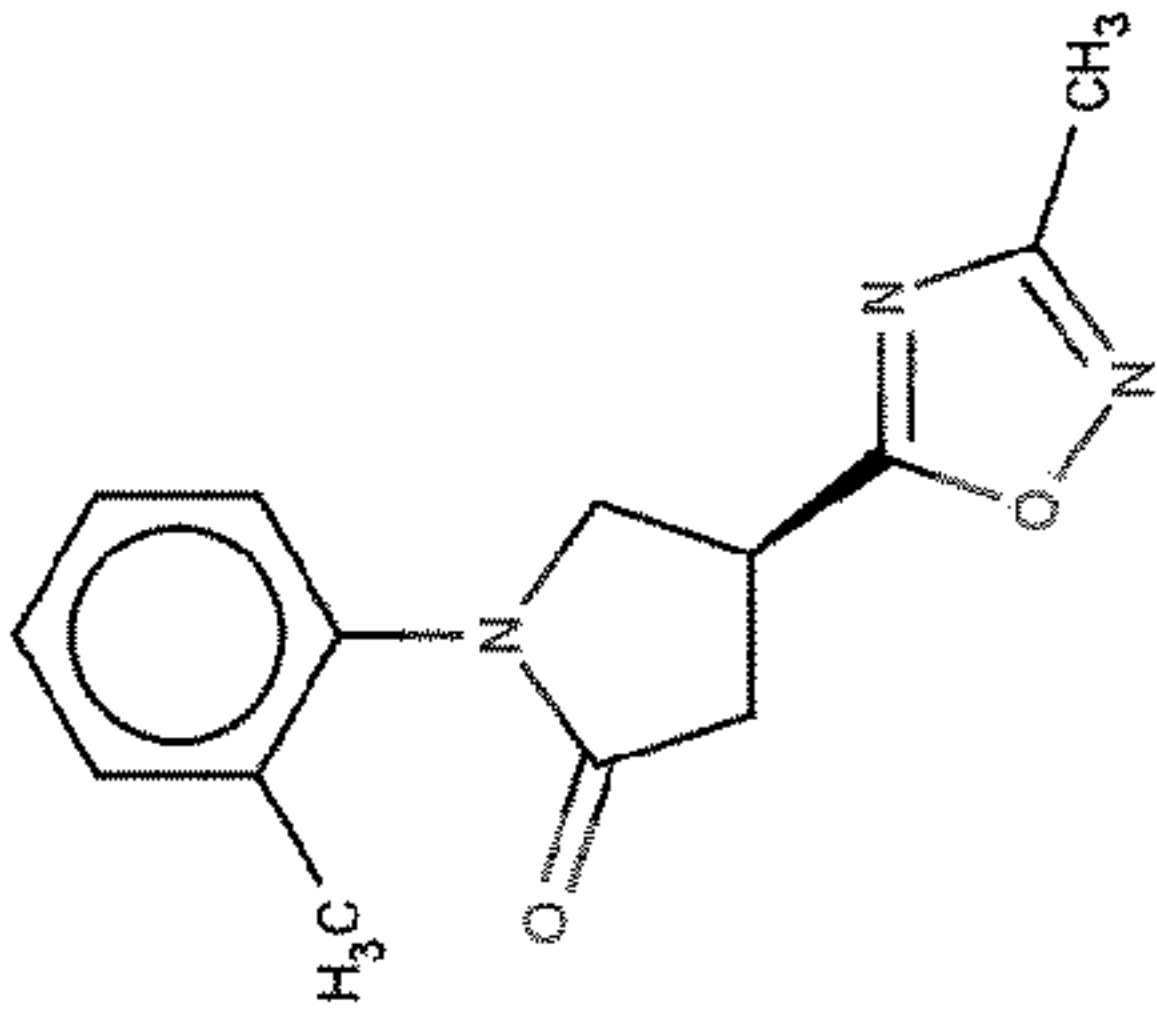
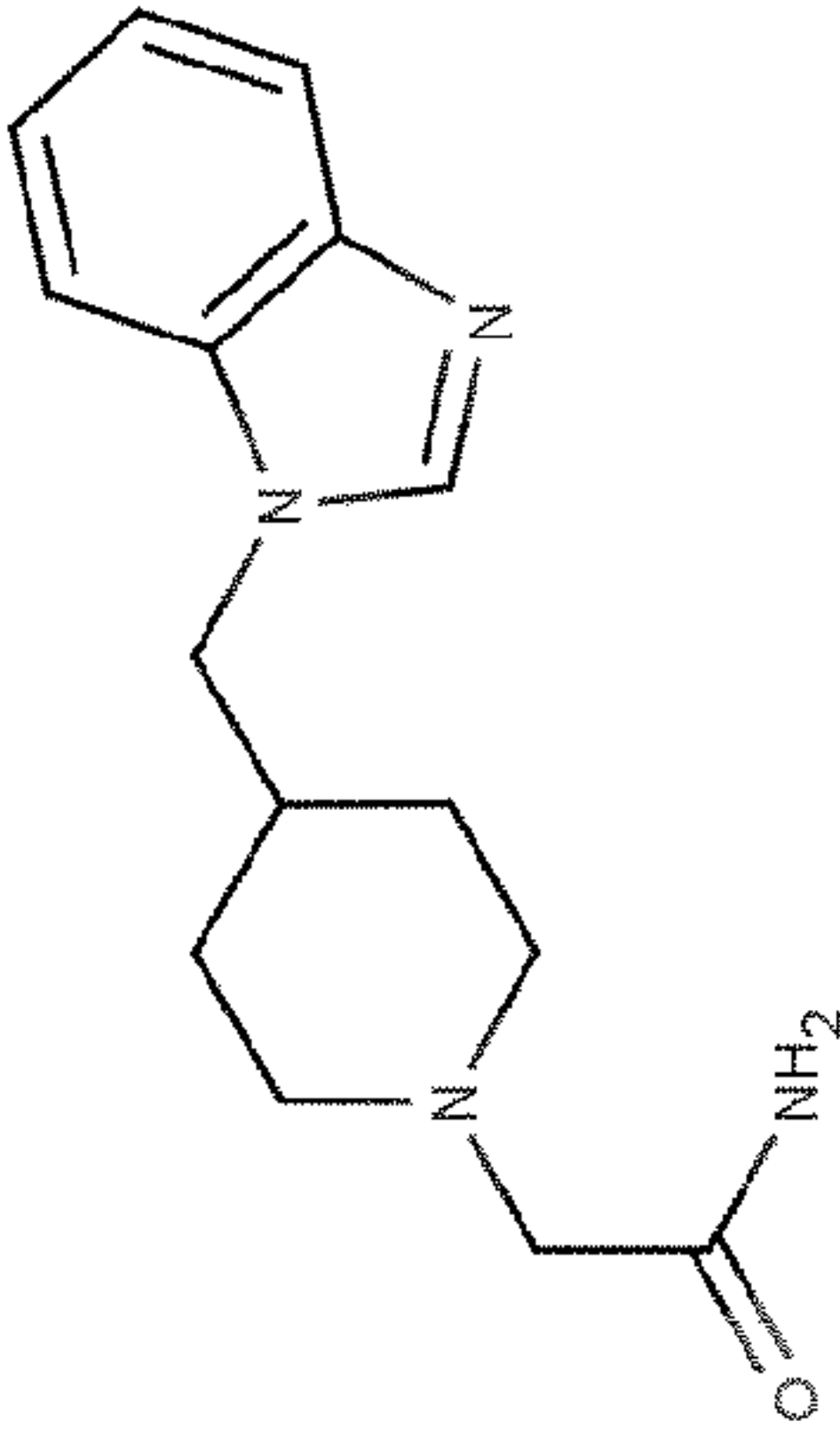
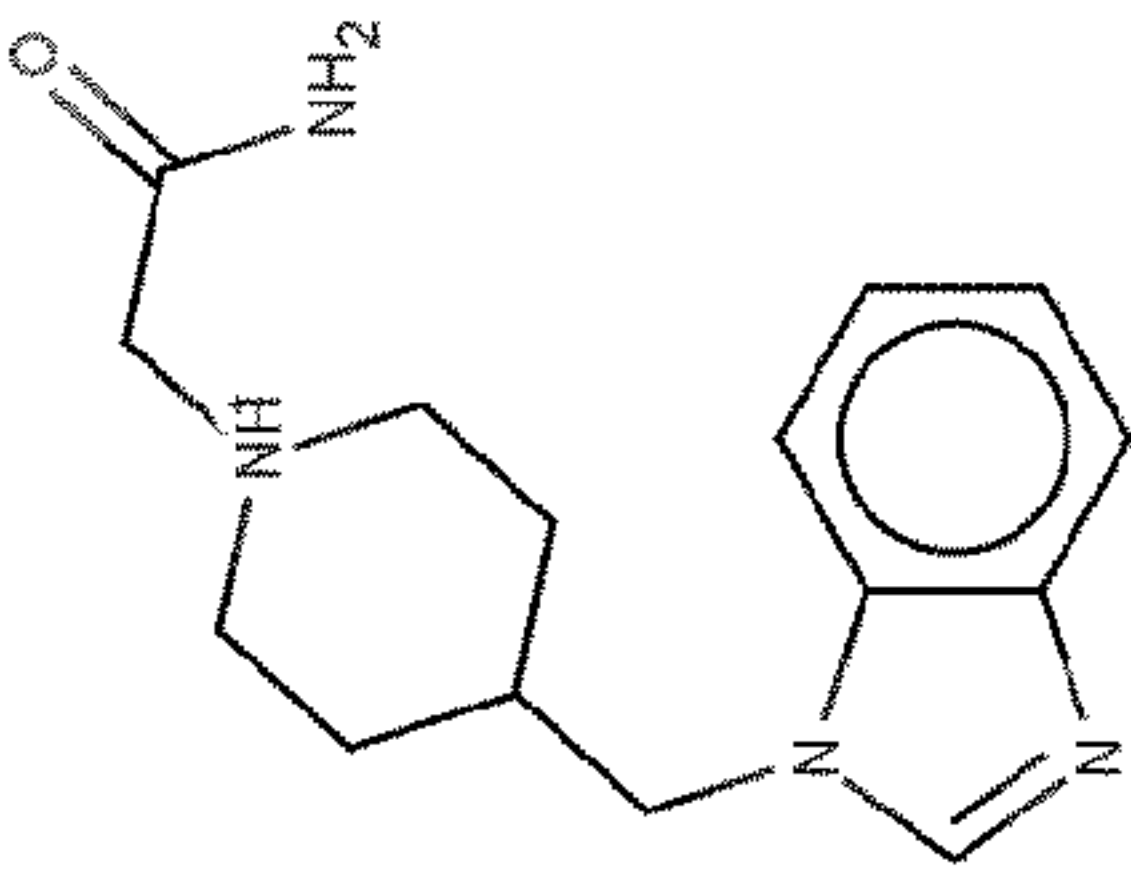
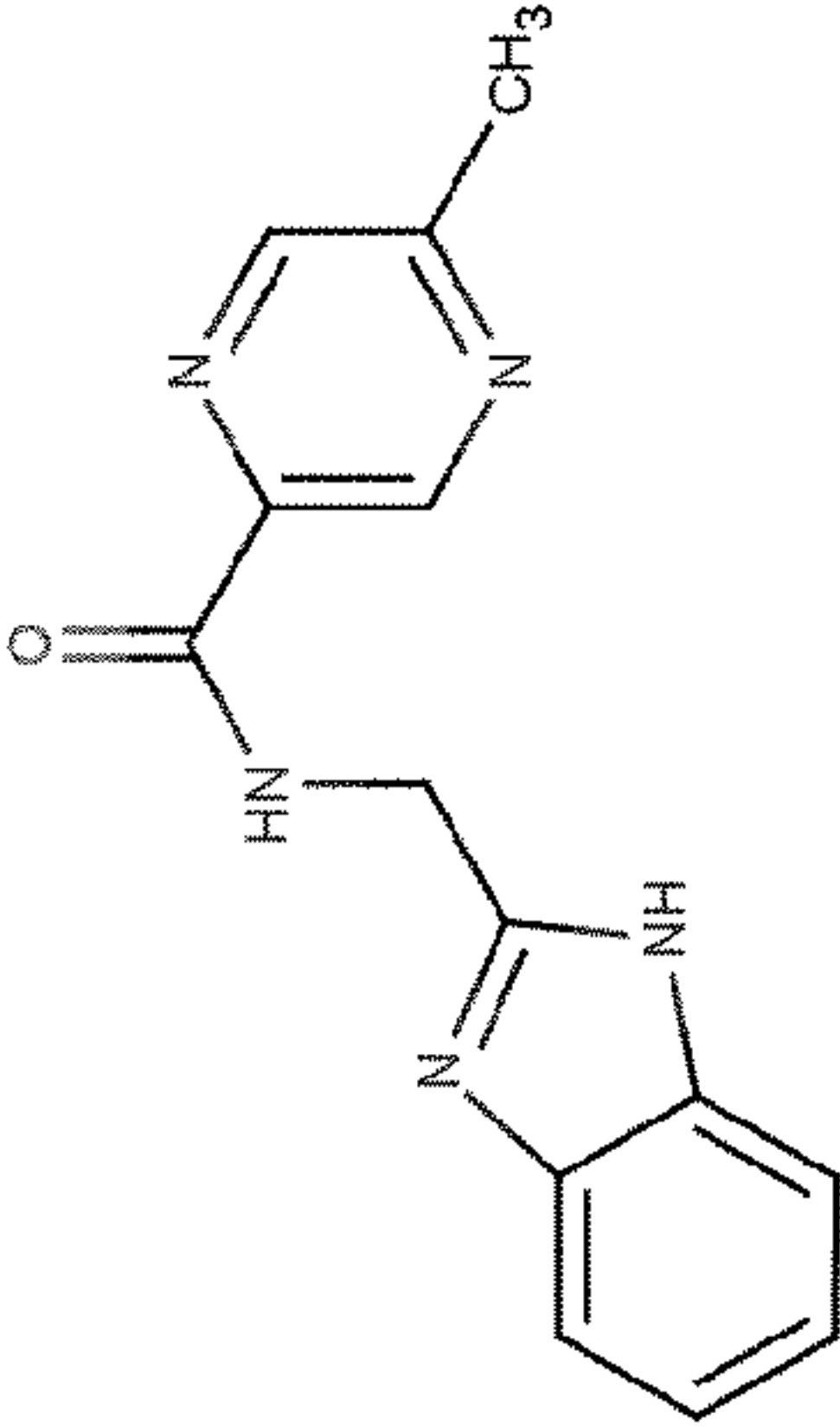
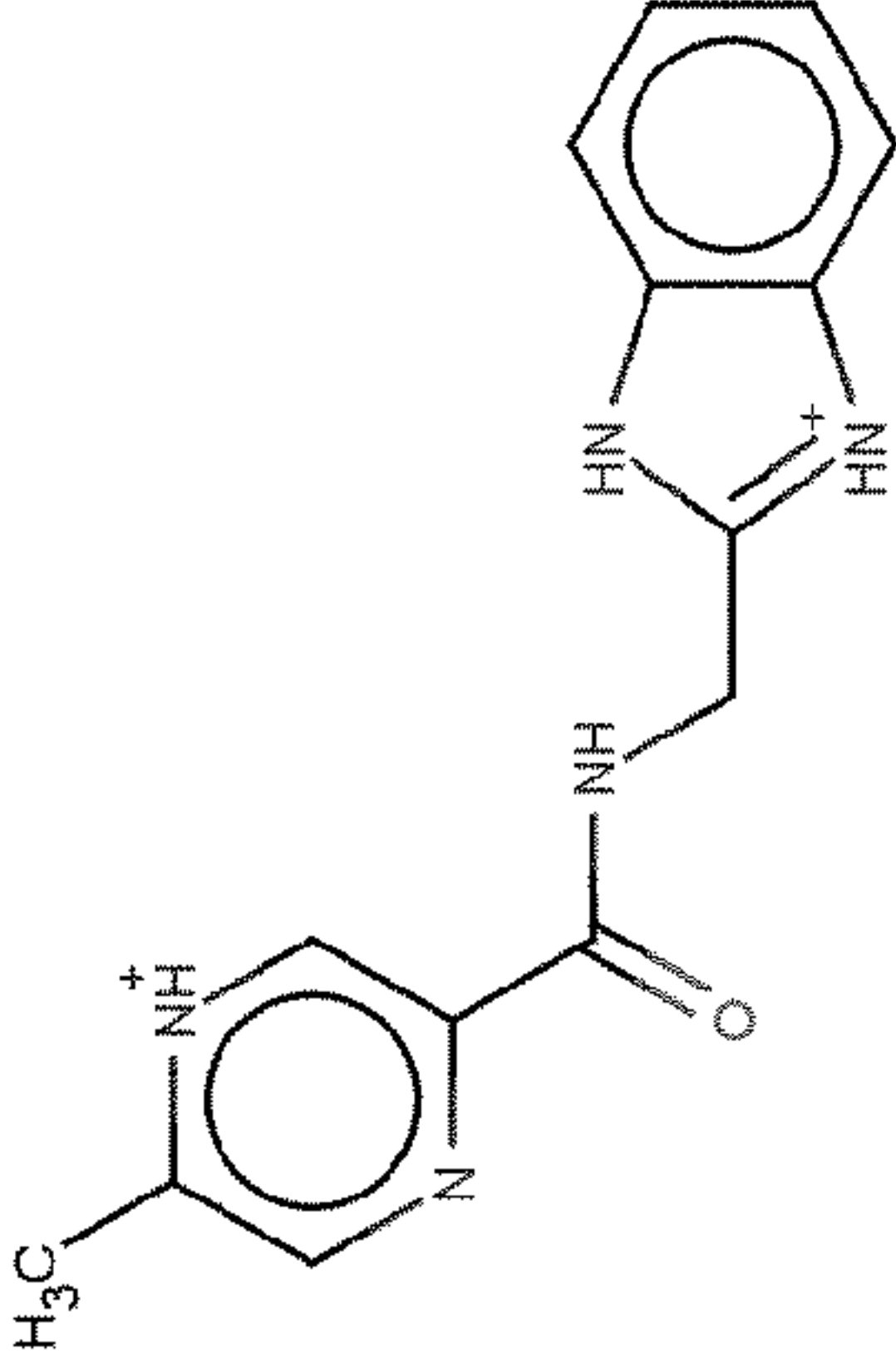
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 211		F3358-0389		F3358-0389_1_1 (1 of 8)
Compound 212		F3368-0119		F3368-0119_1_1 (1 of 4)
Compound 213		F3394-1172		F3394-1172_1_1 (1 of 5)

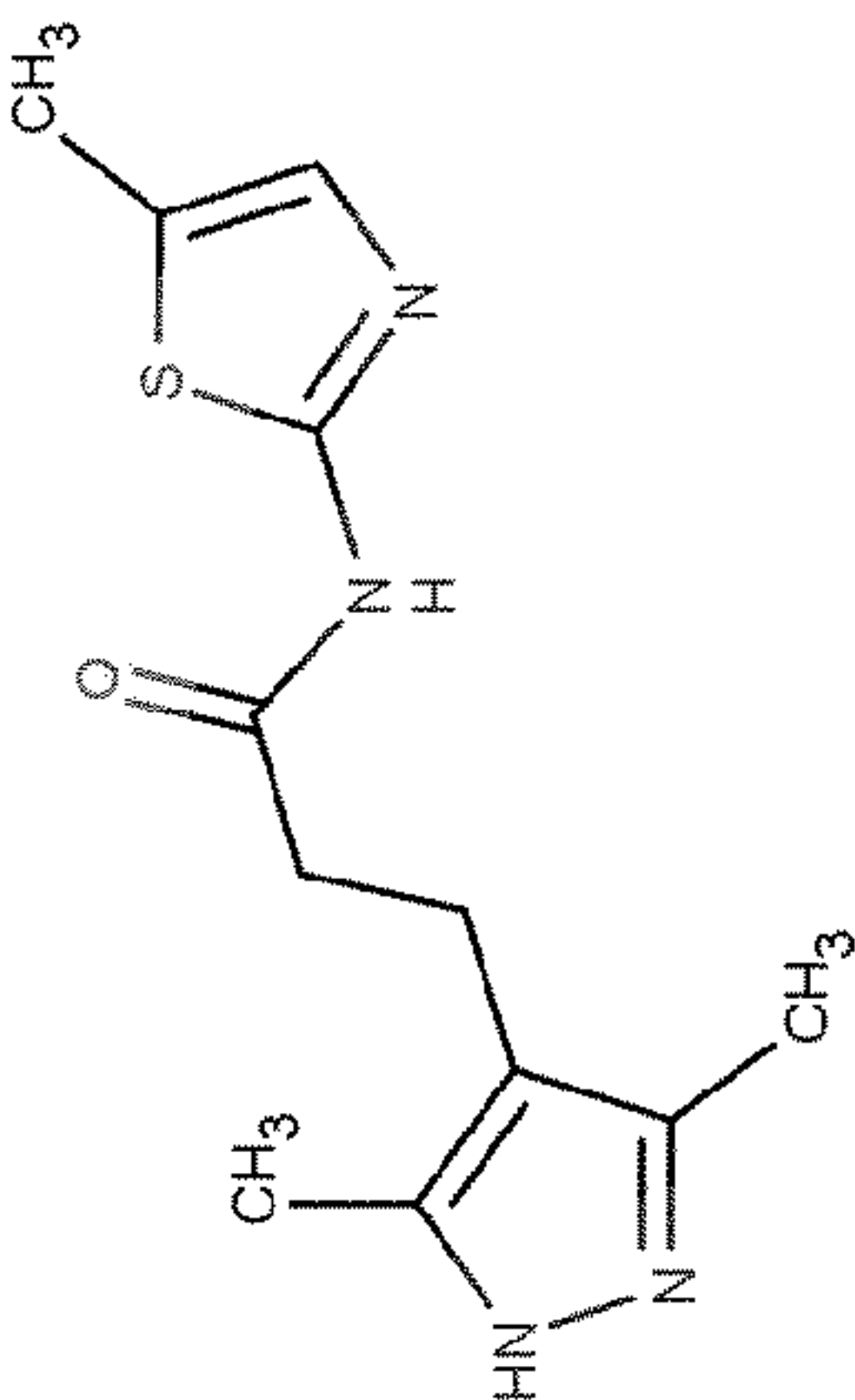
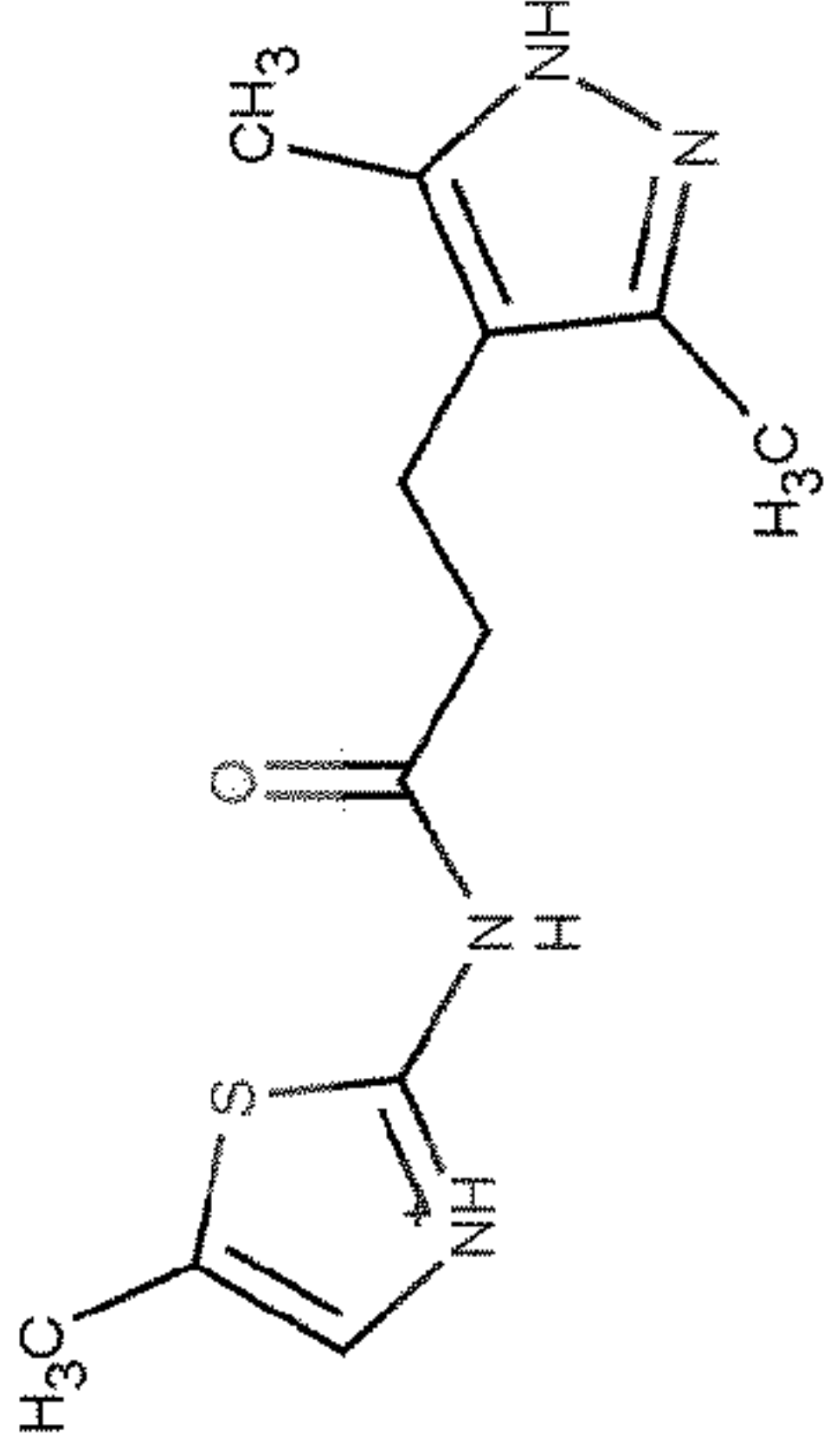
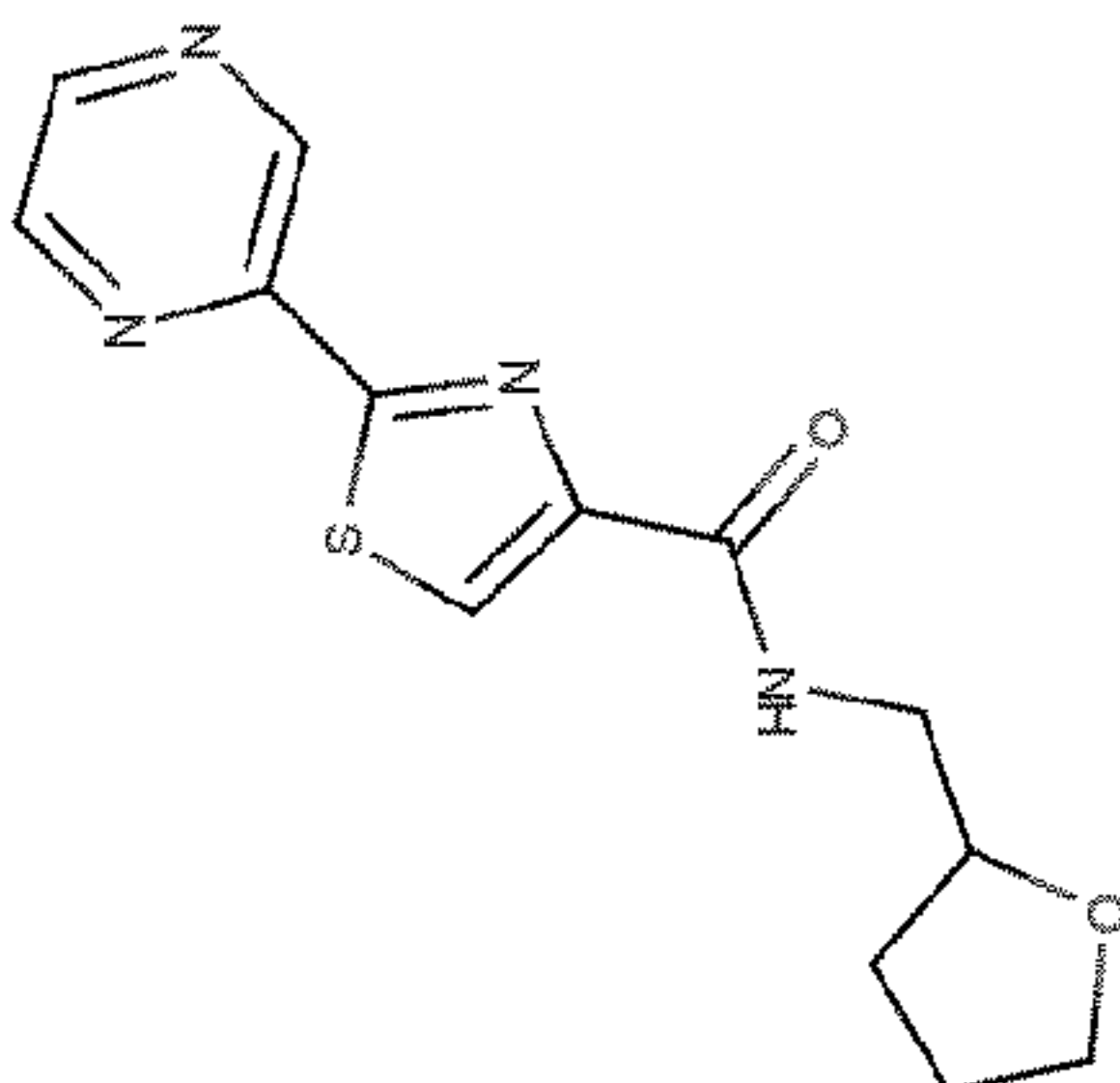
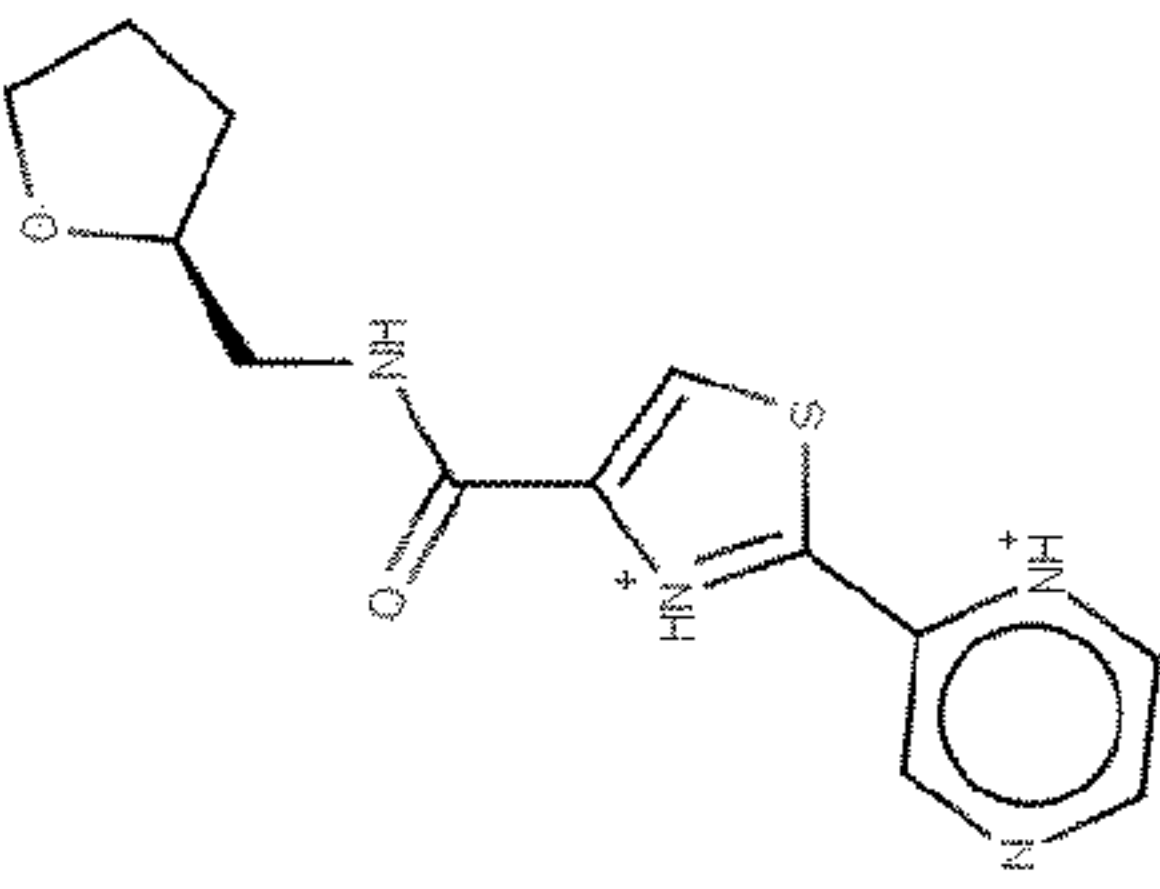
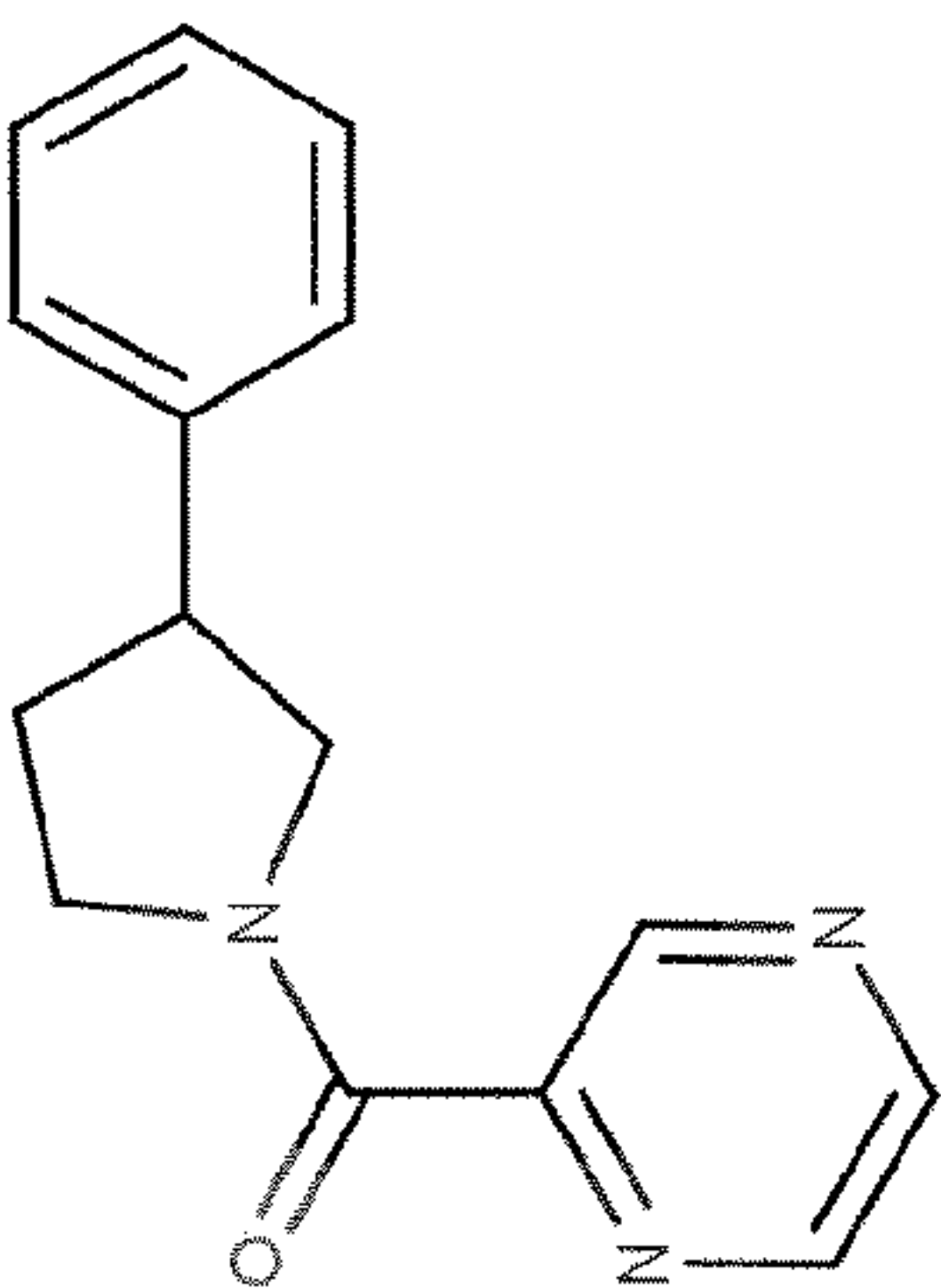
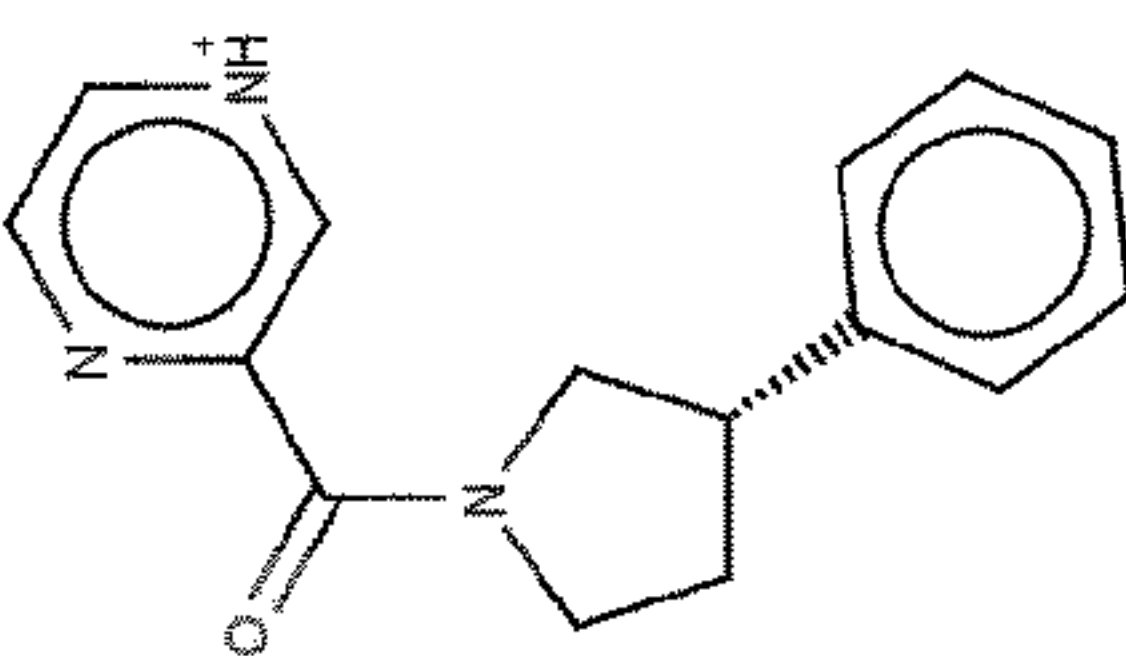
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 214		F3394-1274		F3394-1274_4_1 (1 of 5)
Compound 215		F3406-8965		F3406-8965_39_2 (1 of 158)
Compound 216		F3407-5137		F3407-5137_3_1 (1 of 3)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 217		F5017-0030		F5017-0030_1 (1 of 2)
Compound 218		F5017-0031		F5017-0031_1_1 (1 of 2)
Compound 219		F5017-0074		F5017-0074_1 (1 of 2)

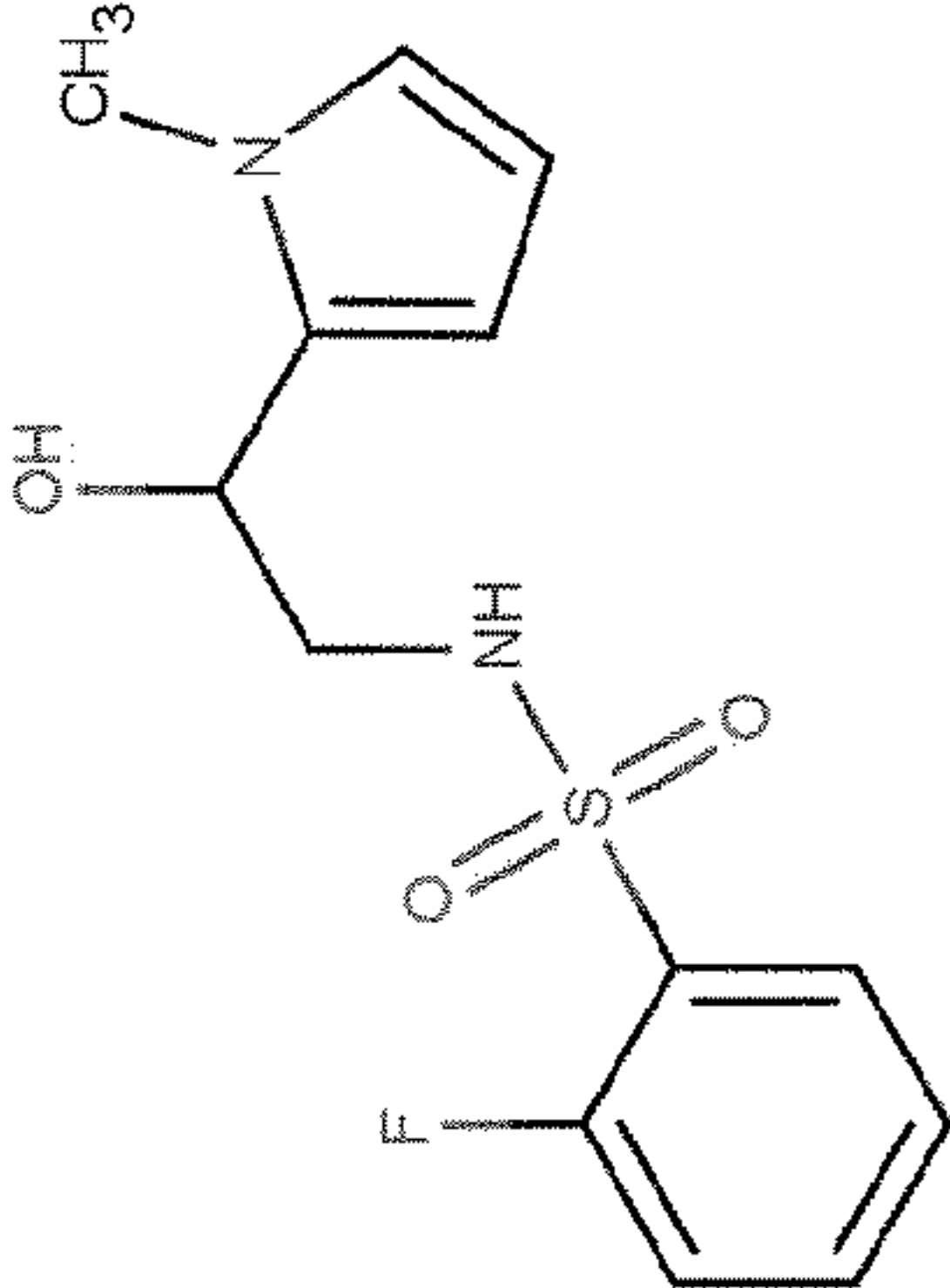
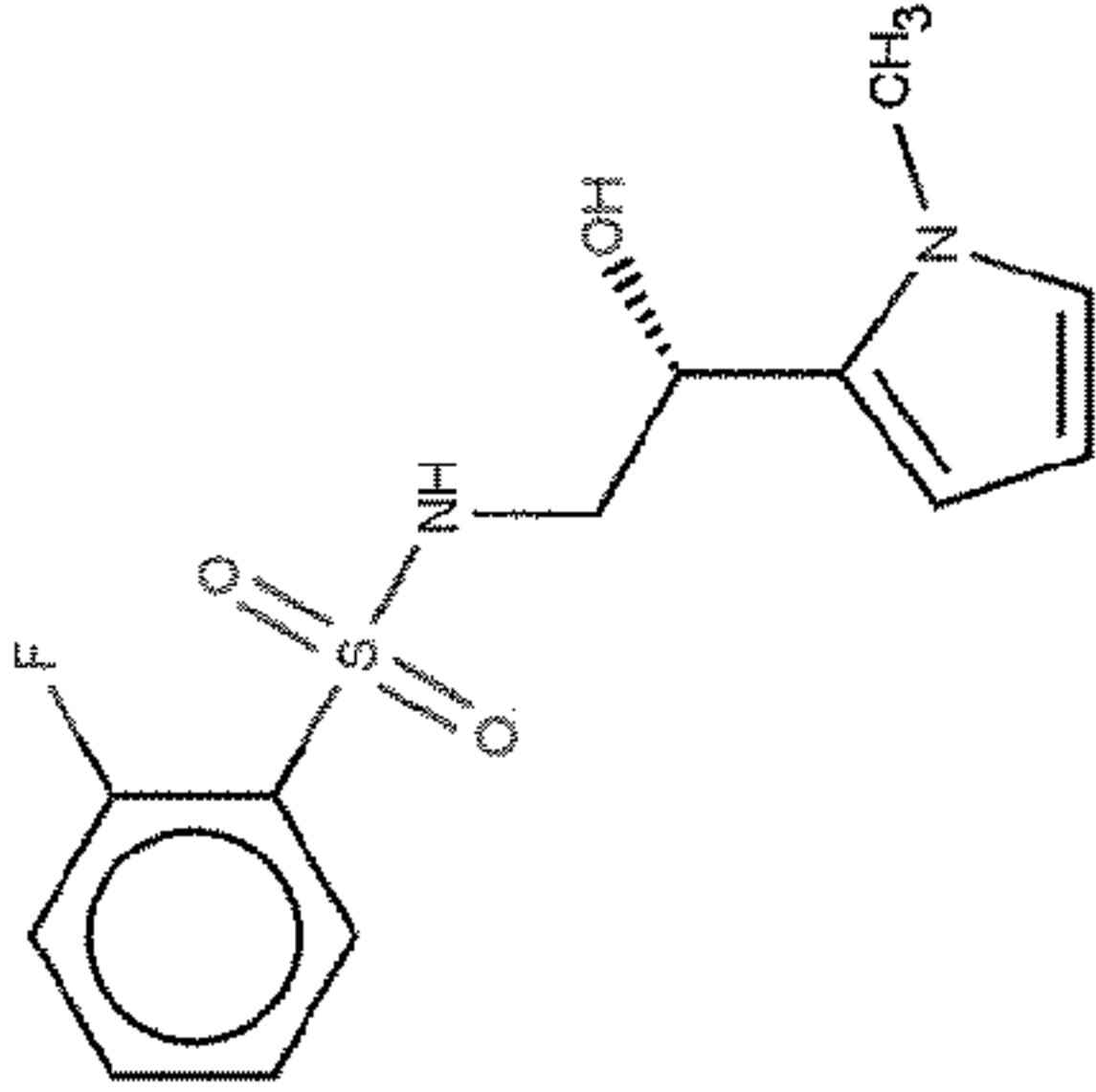
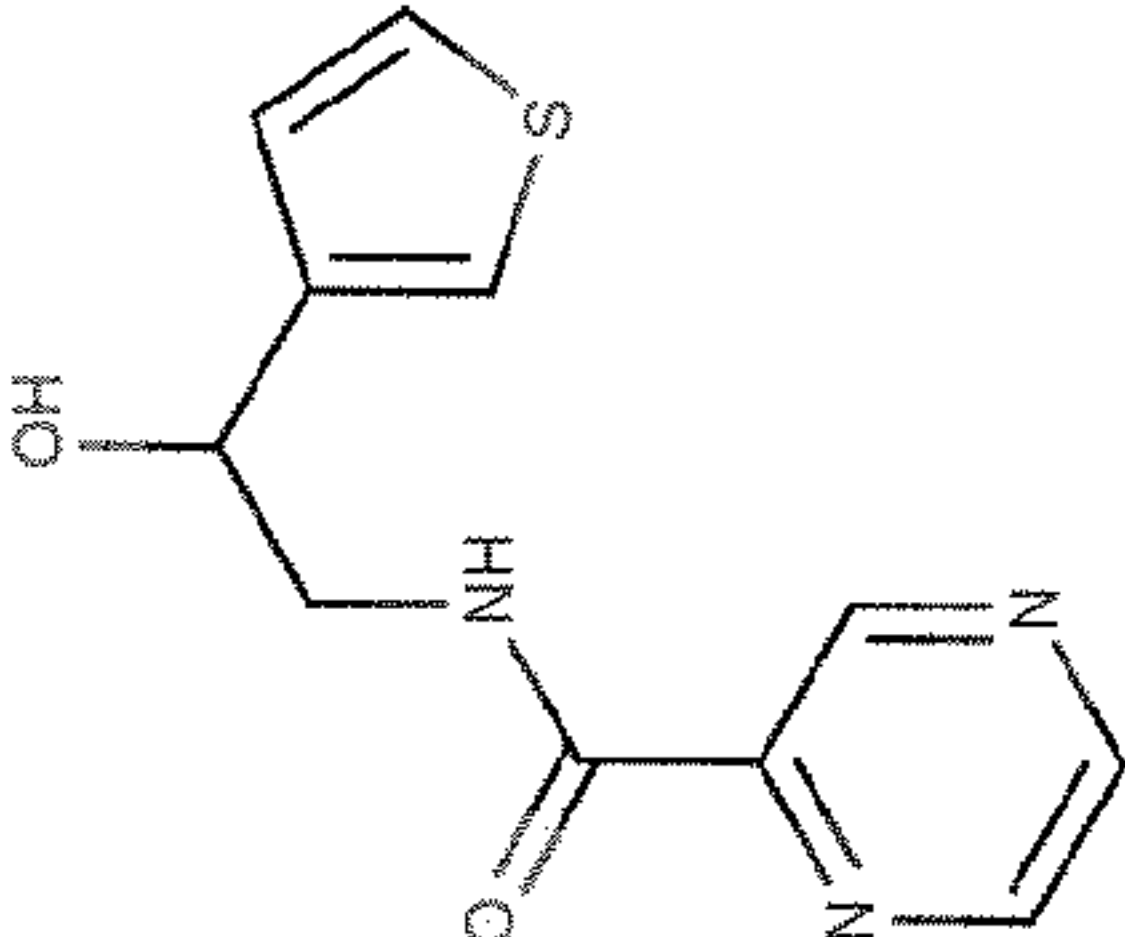
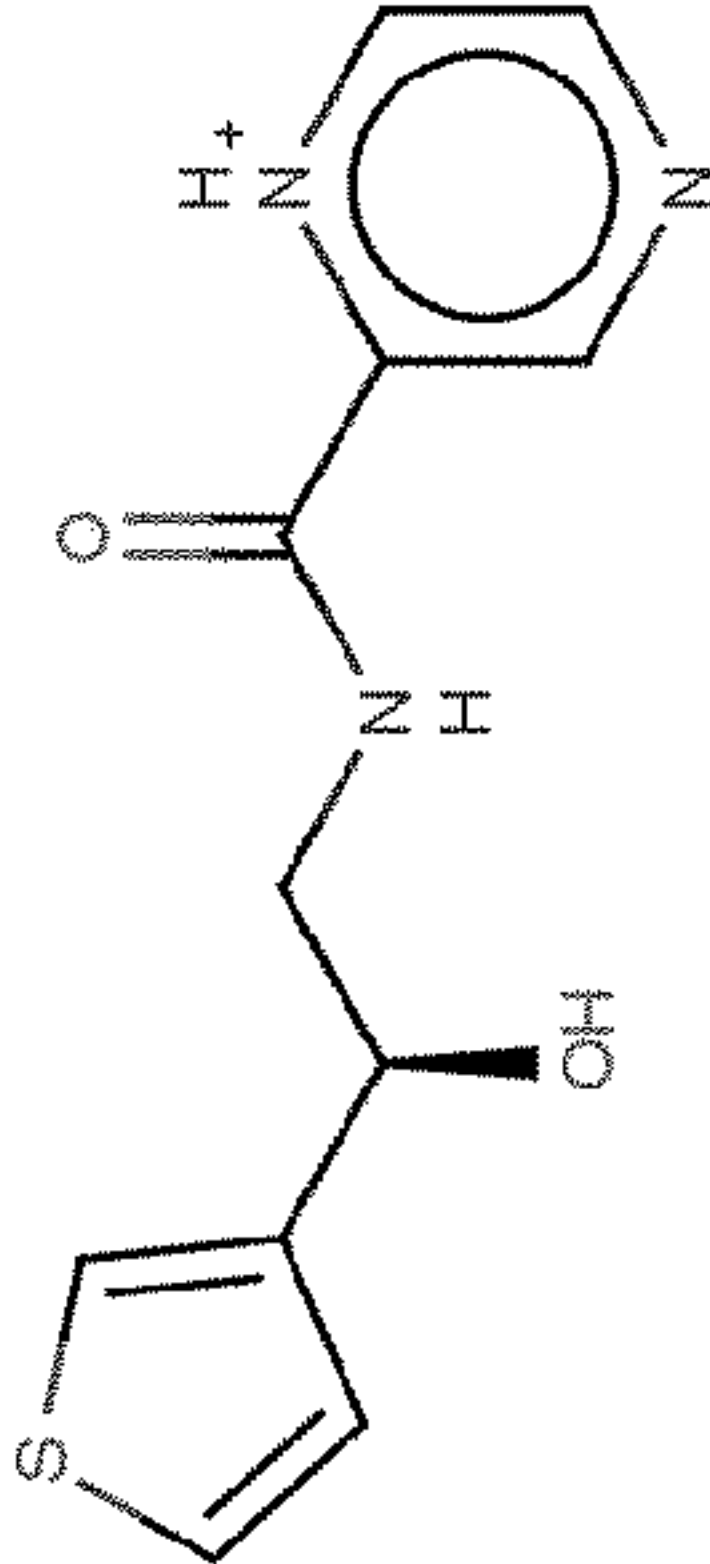
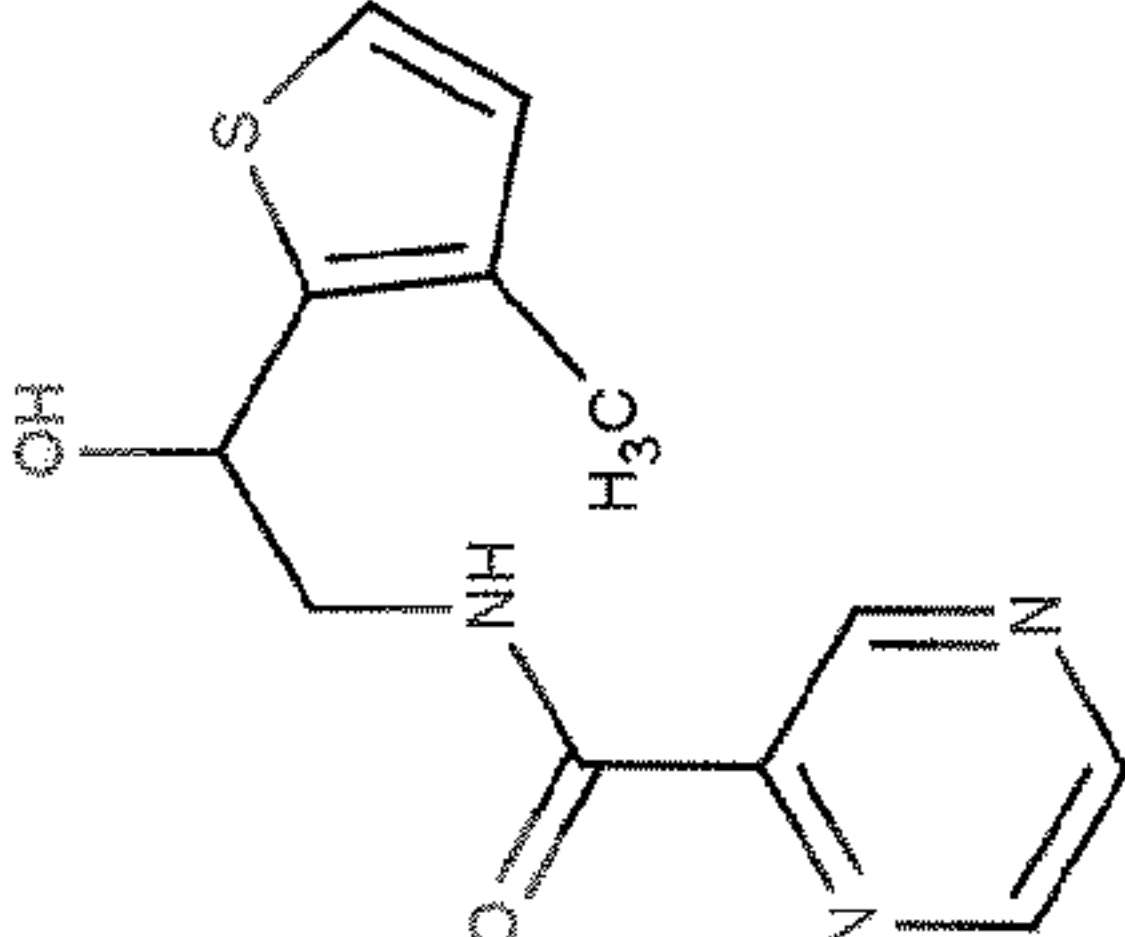
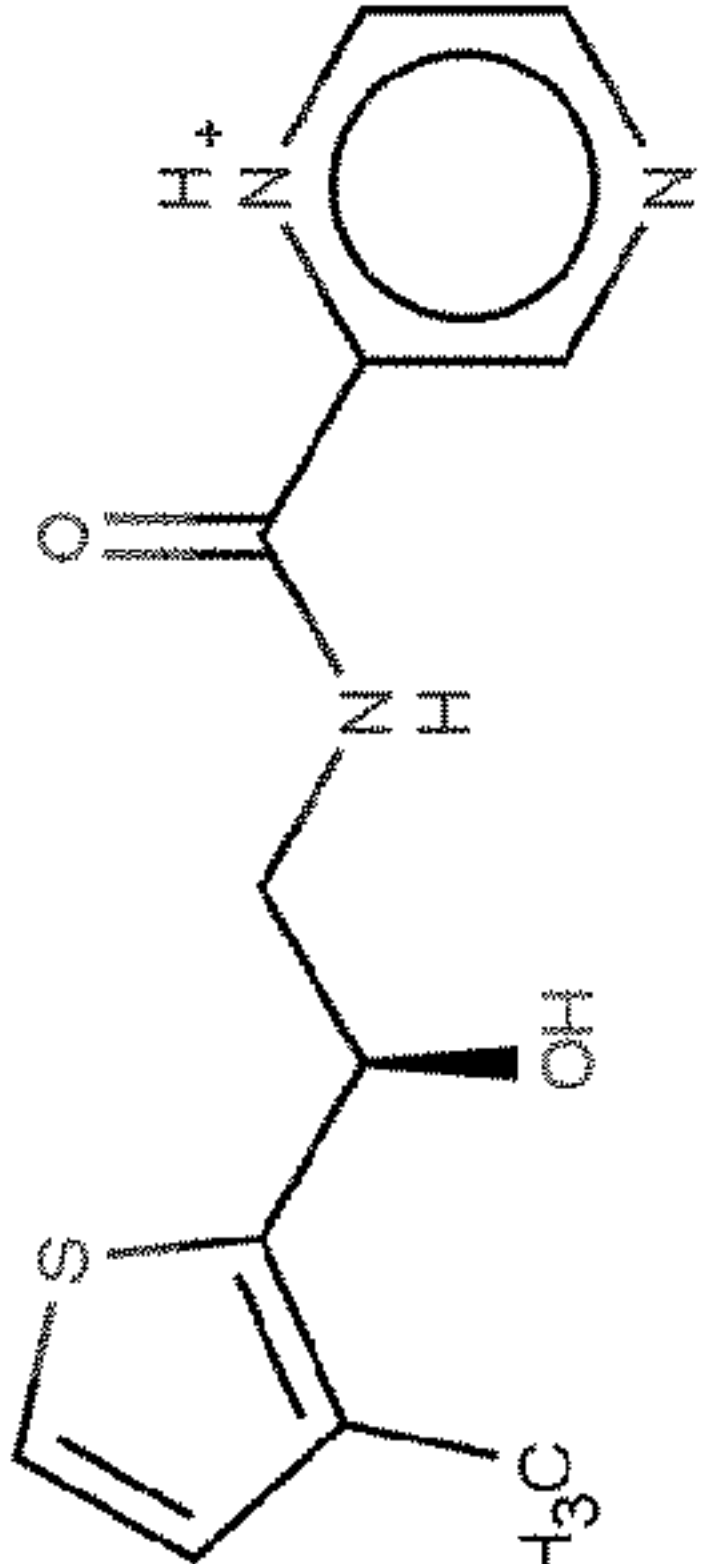
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 220		F5017-0076		F5017-0076_1_1 (1 of 2)
Compound 221		F5080-0092		F5080-0092_14_1 (1 of 4)
Compound 222		F5096-0076		F5096-0076_3_1 (1 of 7)

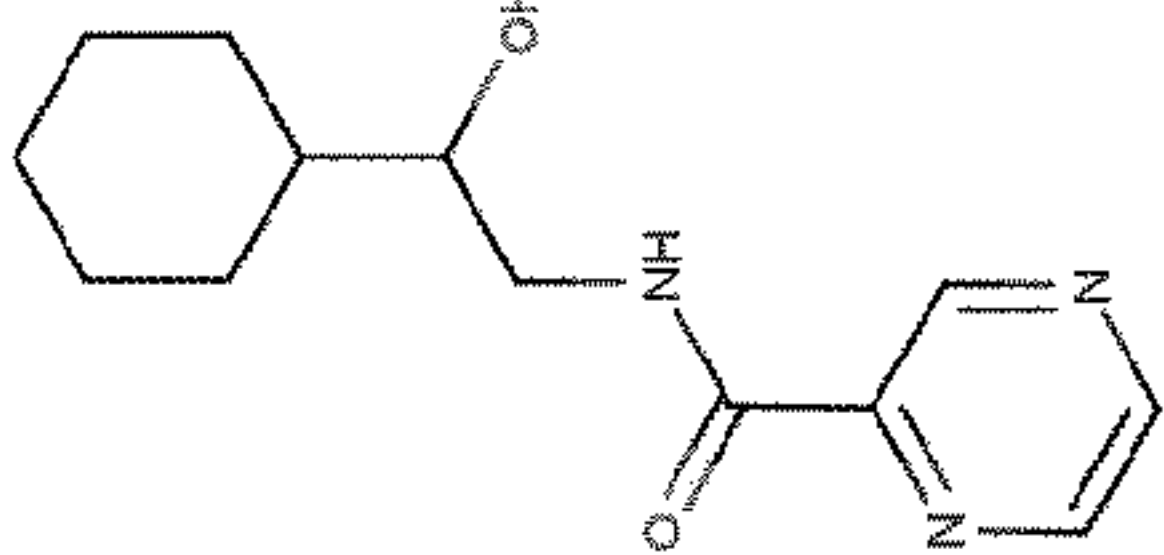
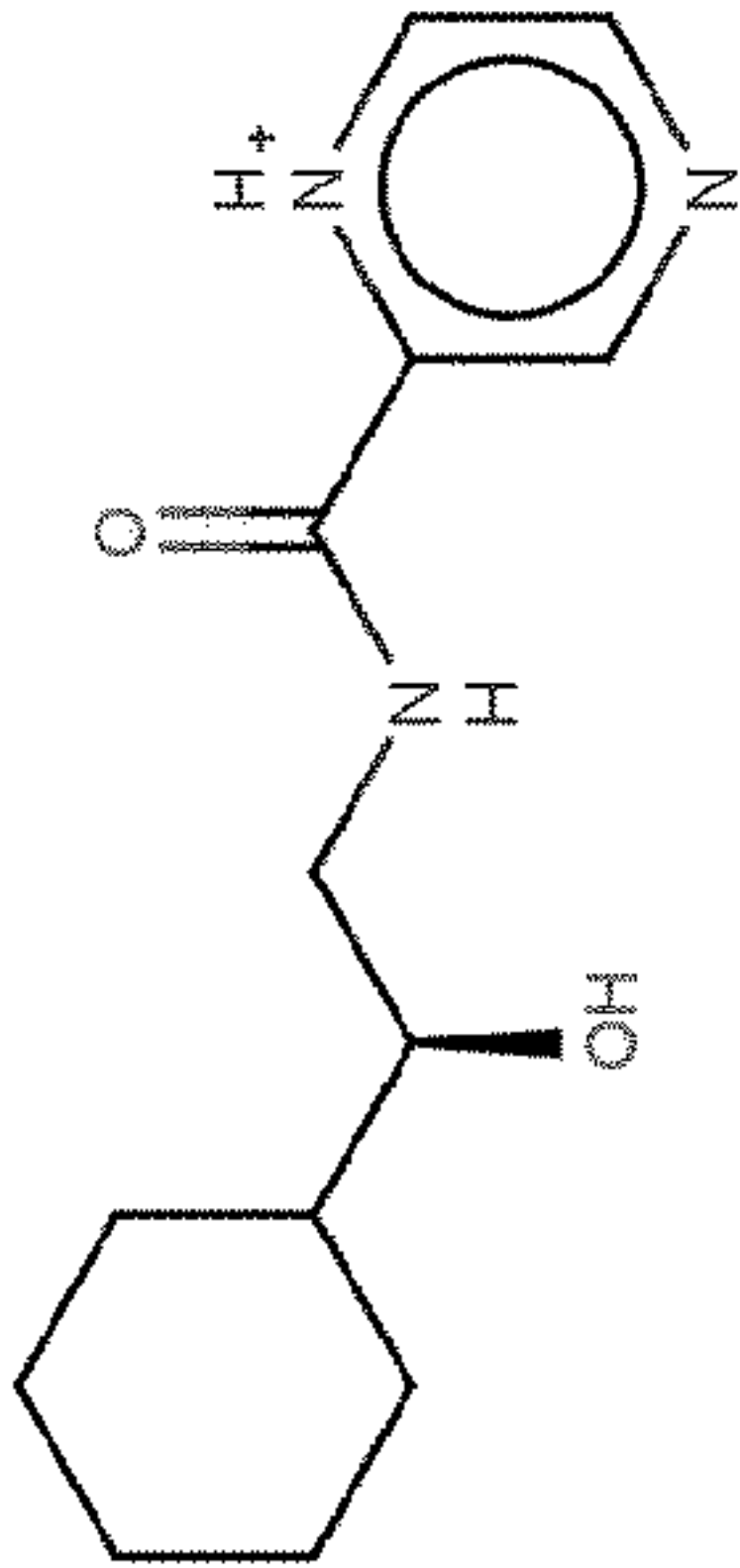
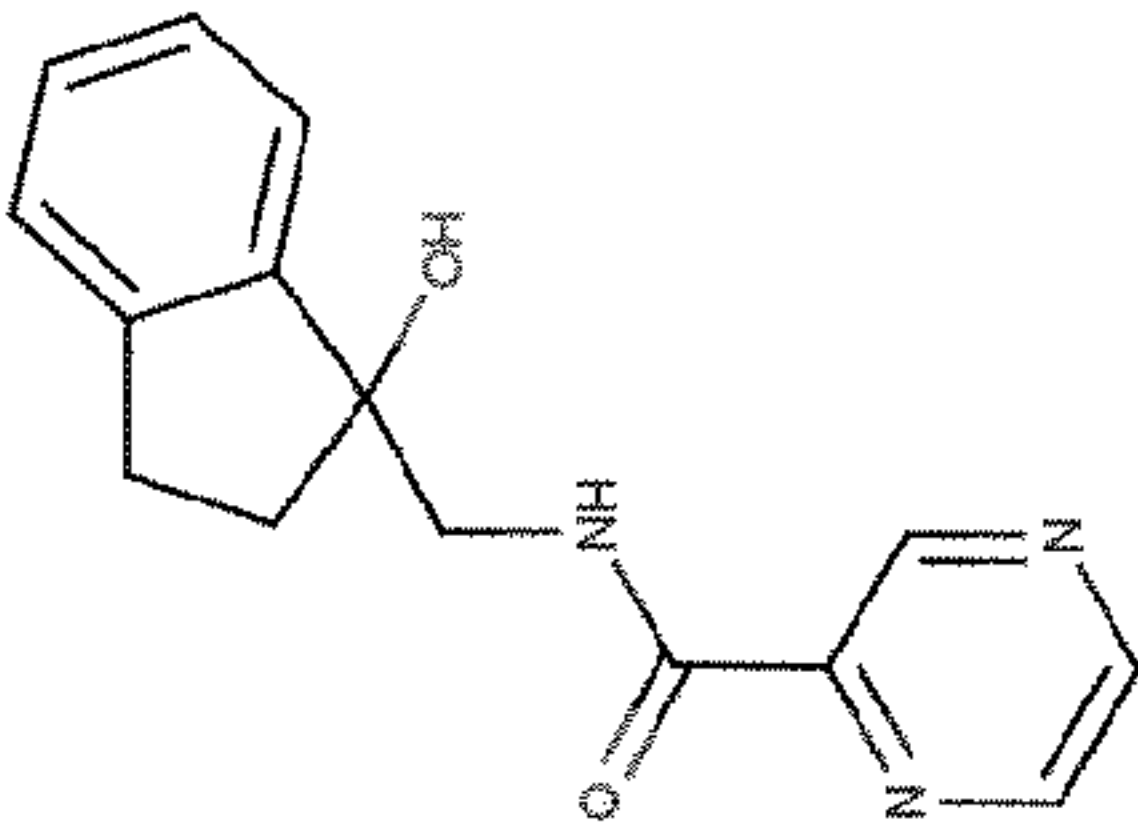
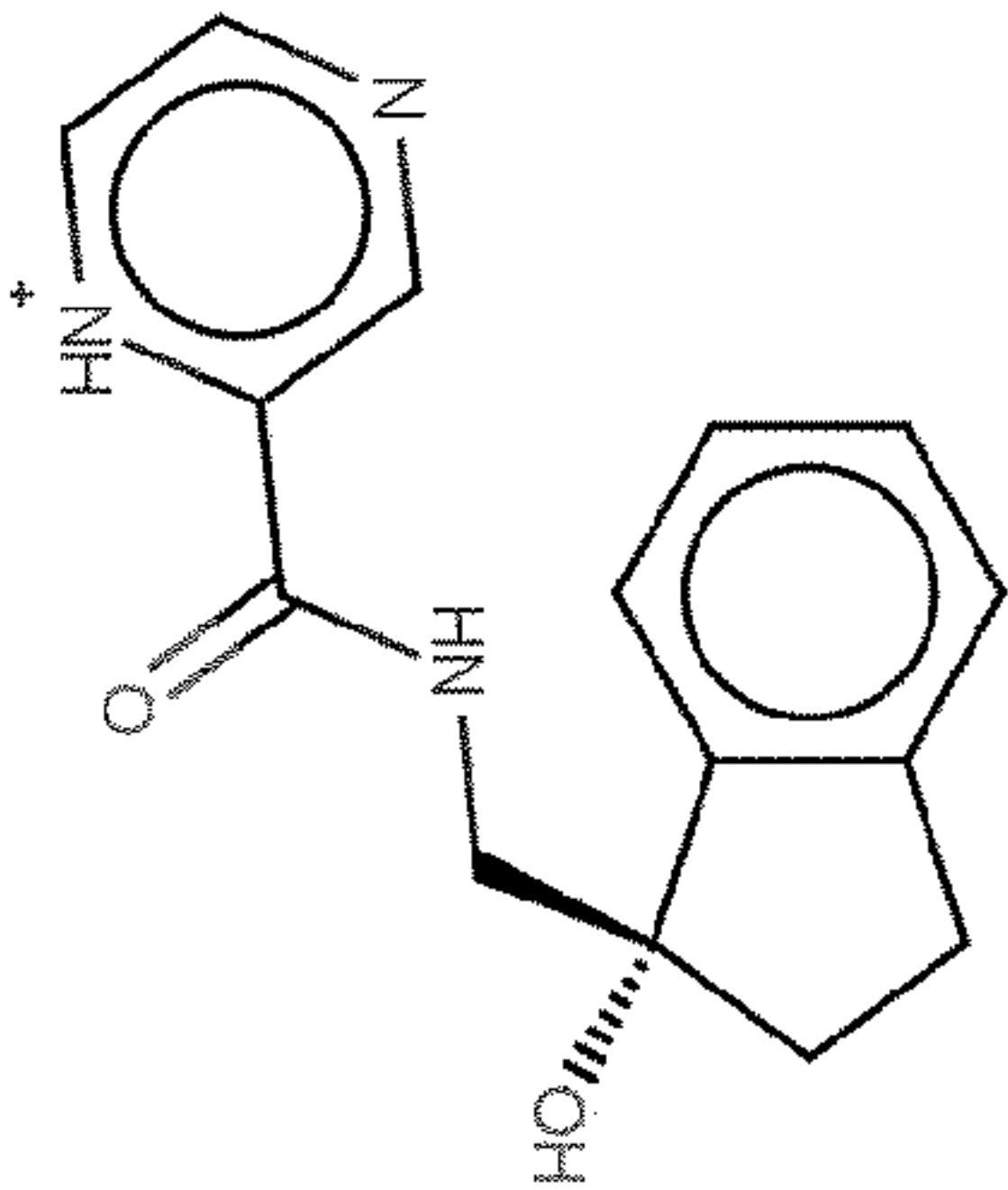
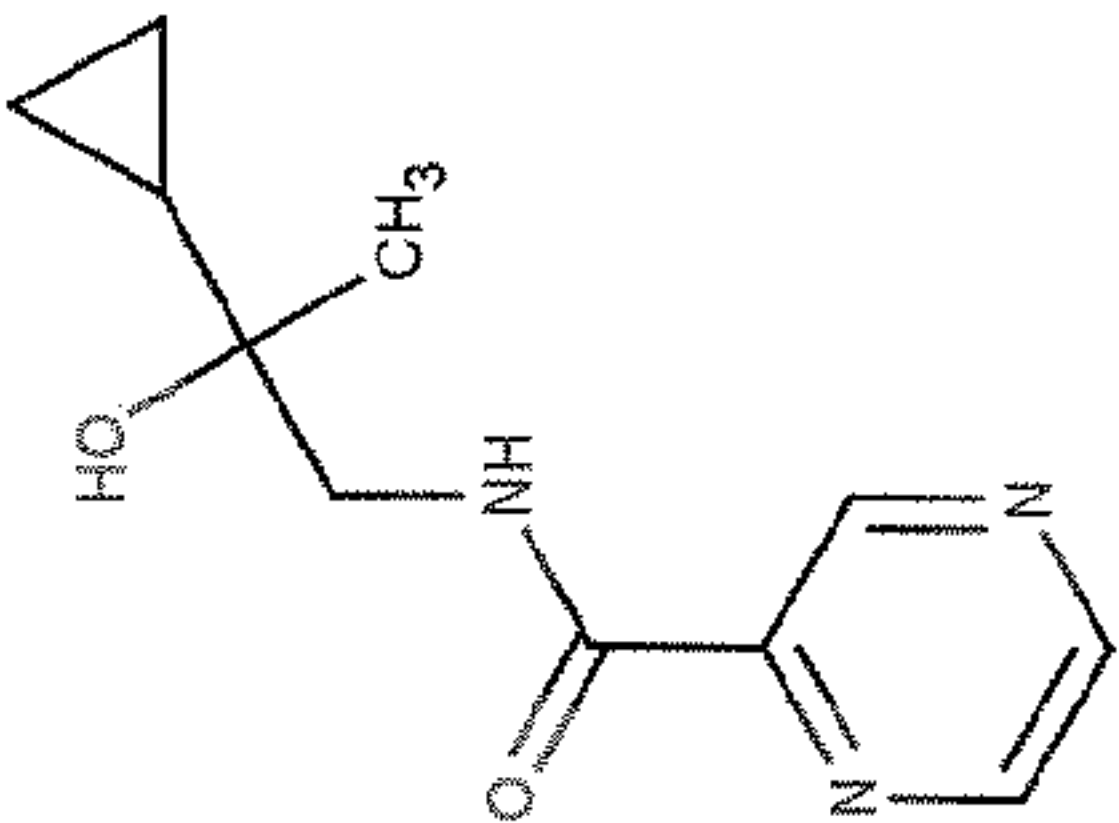
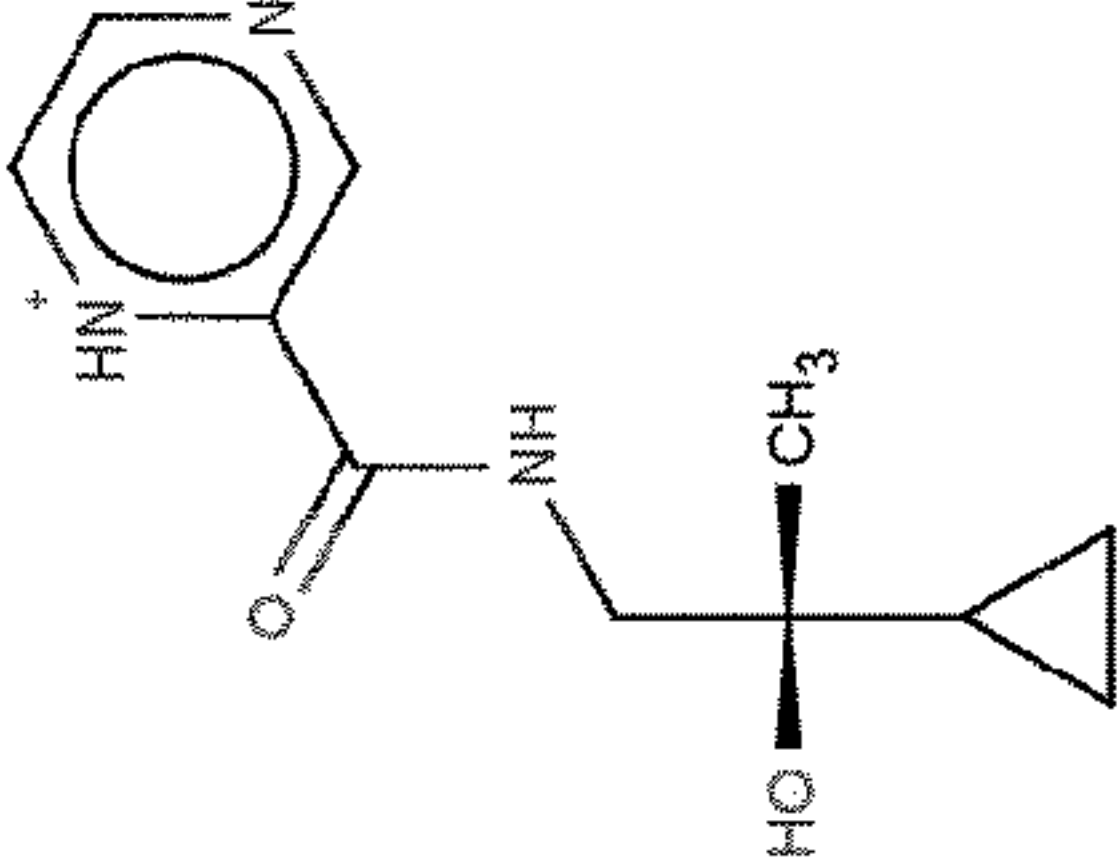
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 223		F5096-0179		F5096-0179_3_1 (1 of 7)
Compound 224		F5302-0092		F5302-0092_1_1 (1 of 2)
Compound 225		F5461-1122		F5461-1122_1_1 (1 of 4)

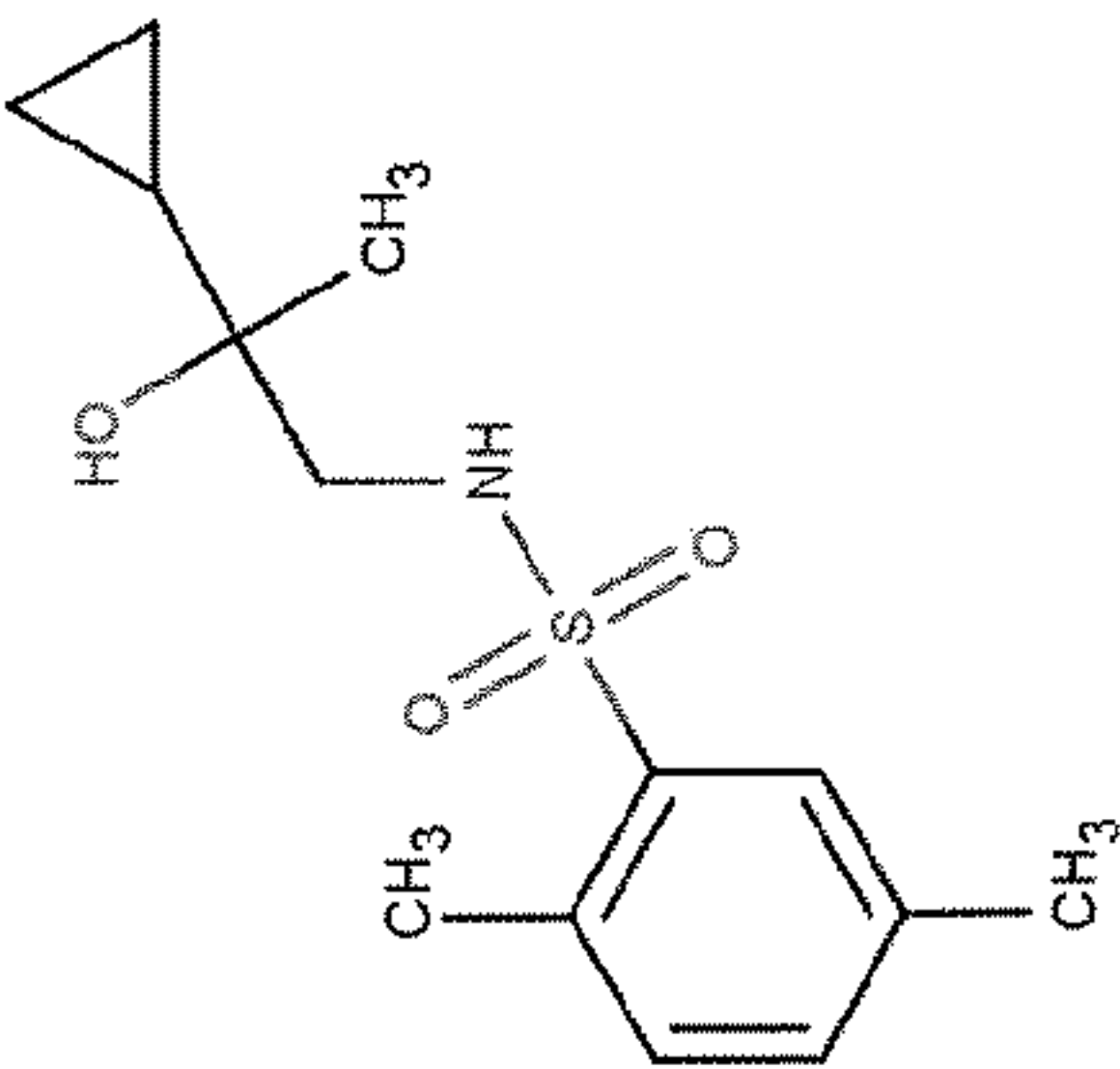
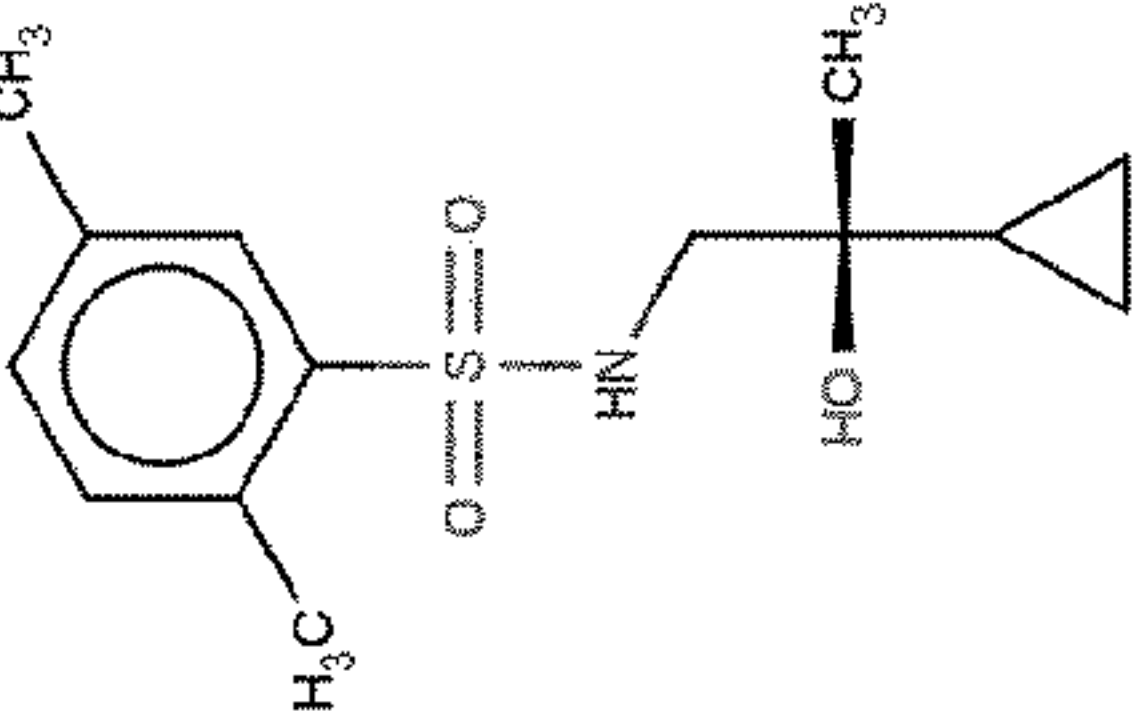
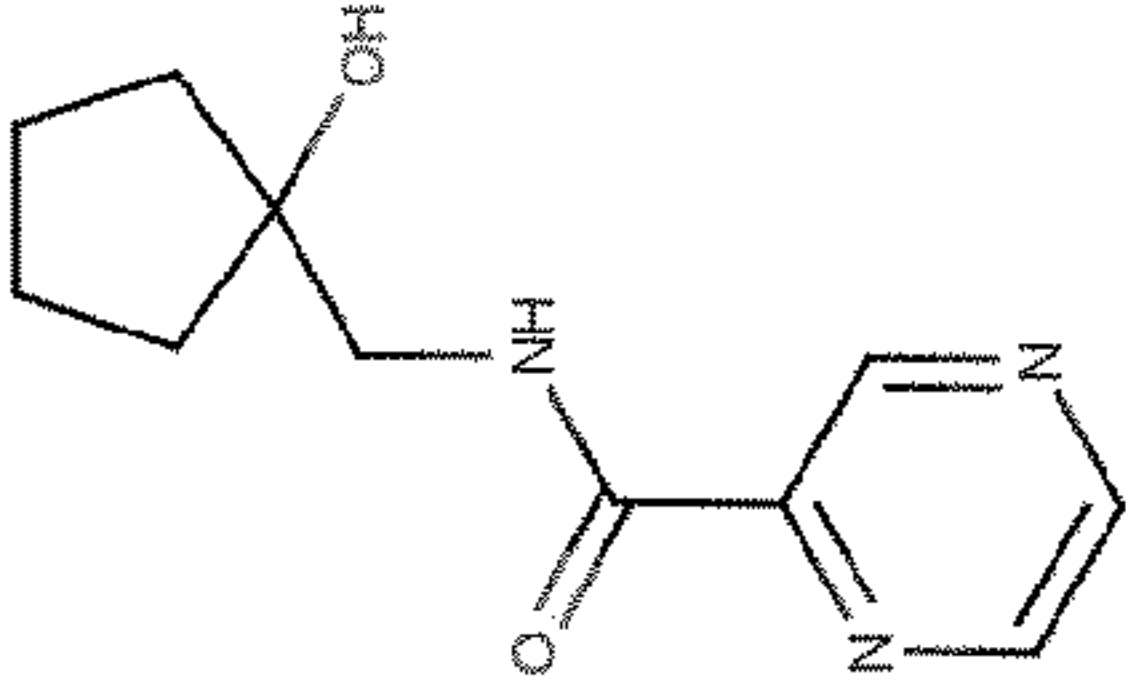
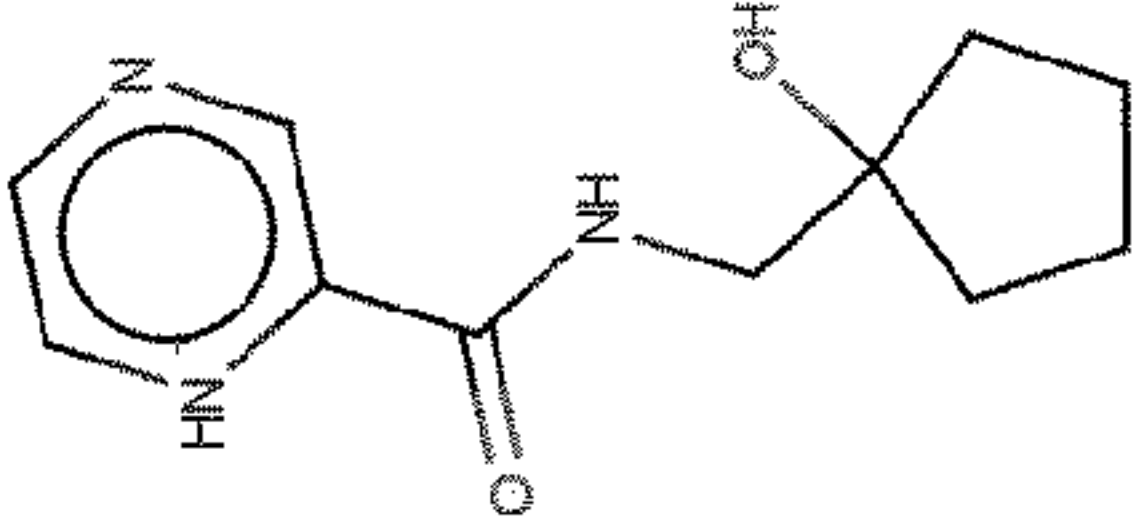
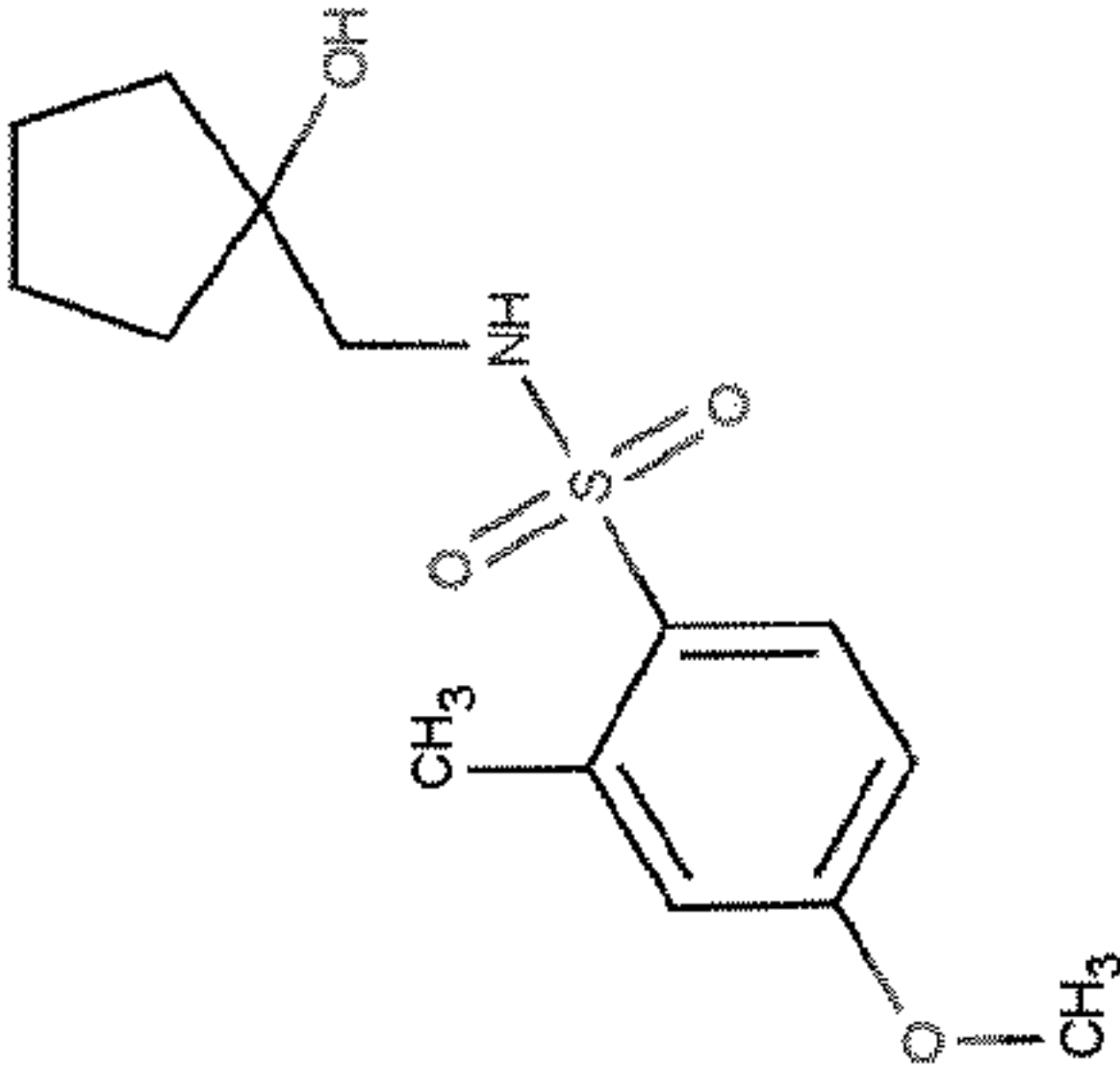
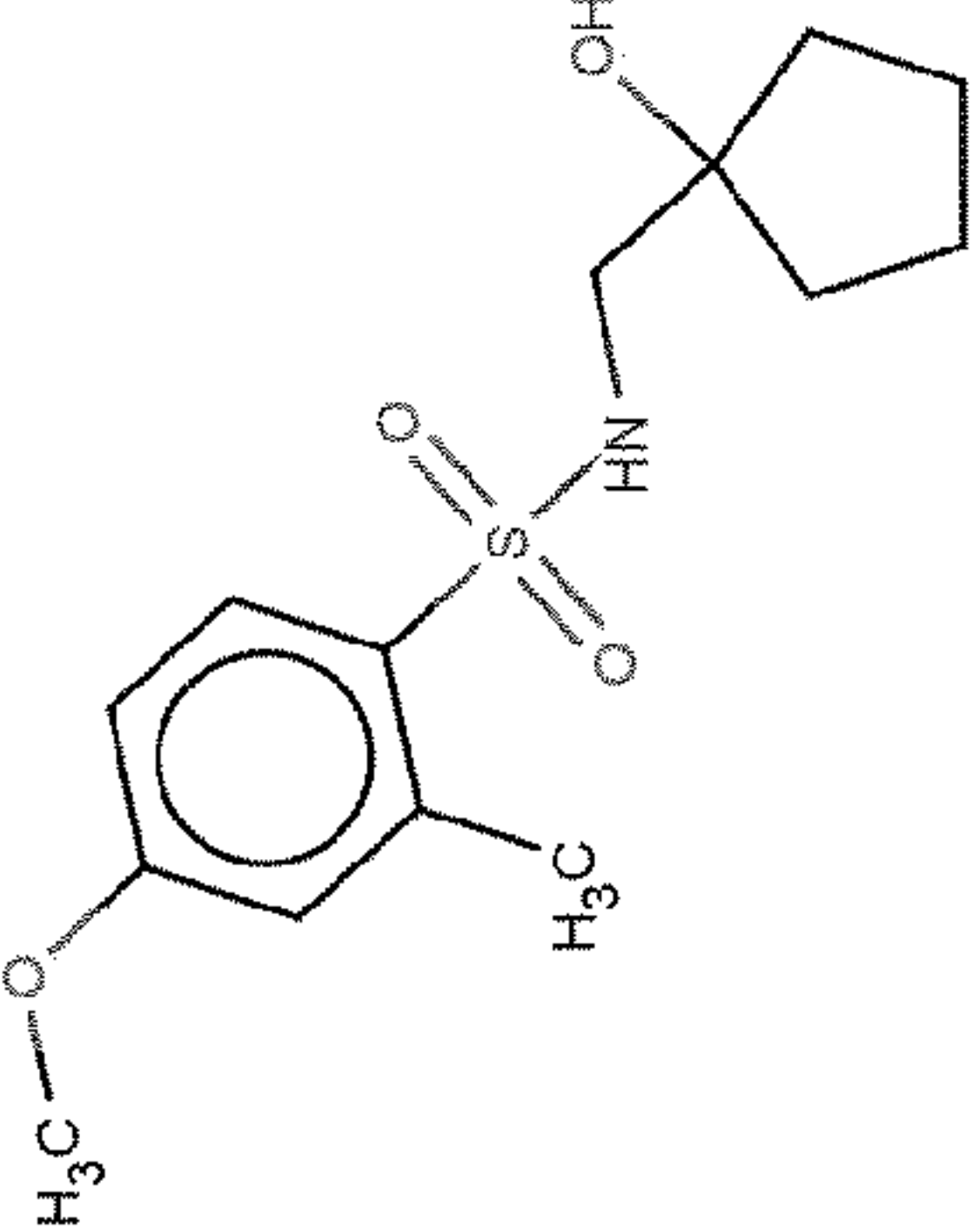
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 226		F5668-0082		F5668-0082_1 (1 of 4)
Compound 227		F5754-0084		F5754-0084_2_1 (1 of 4)
Compound 228		F5791-2428		F5791-2428_5_1 (1 of 6)

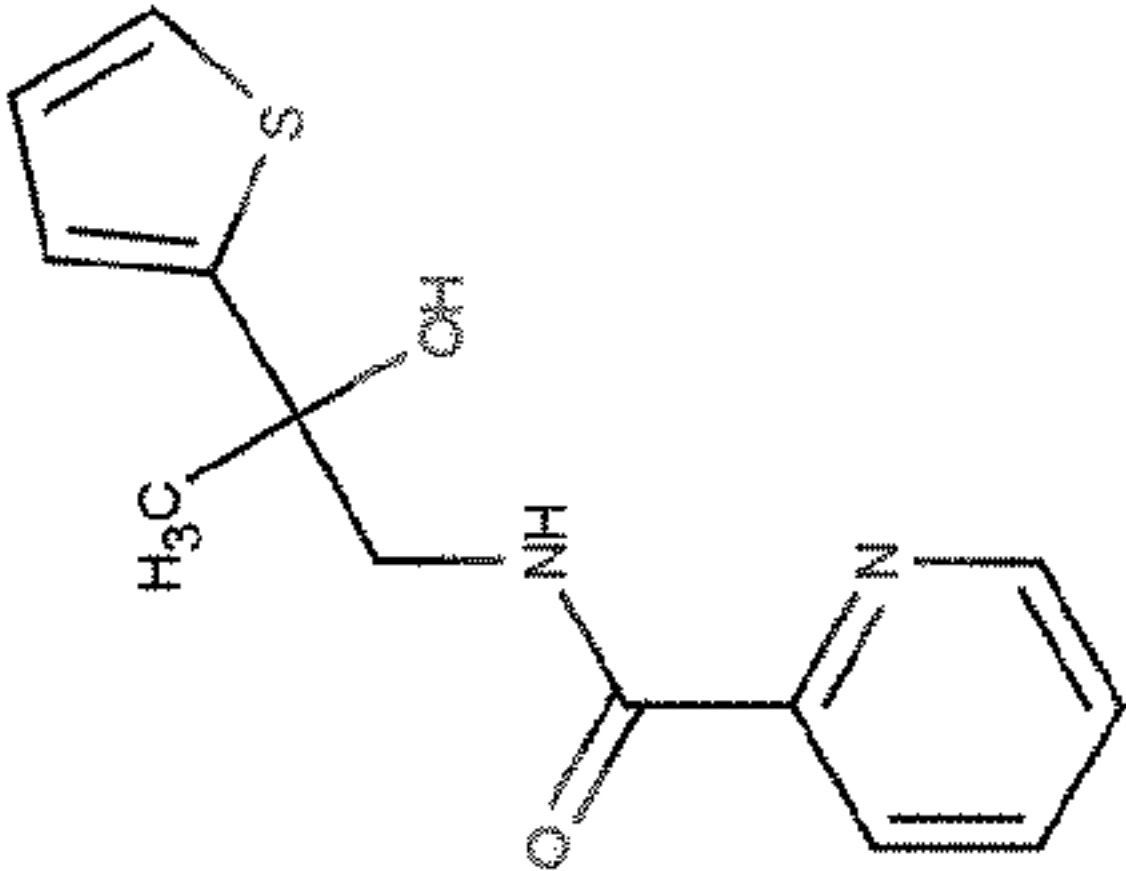
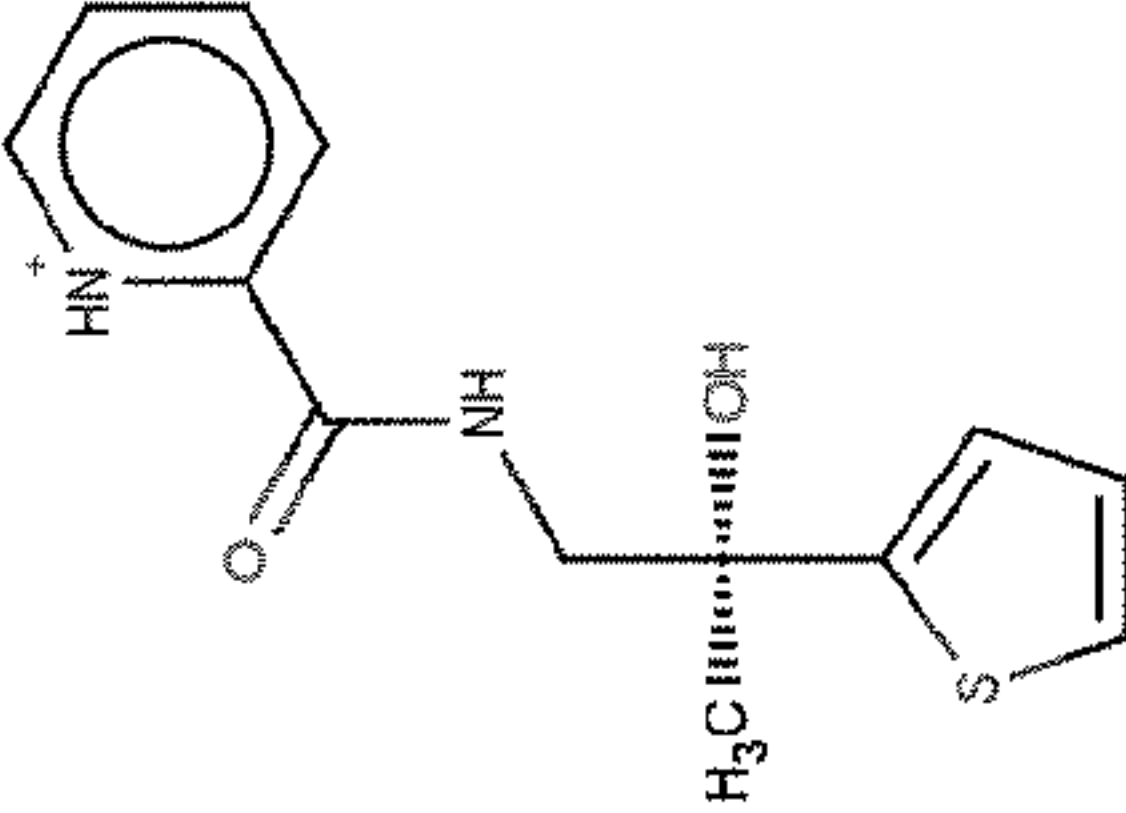
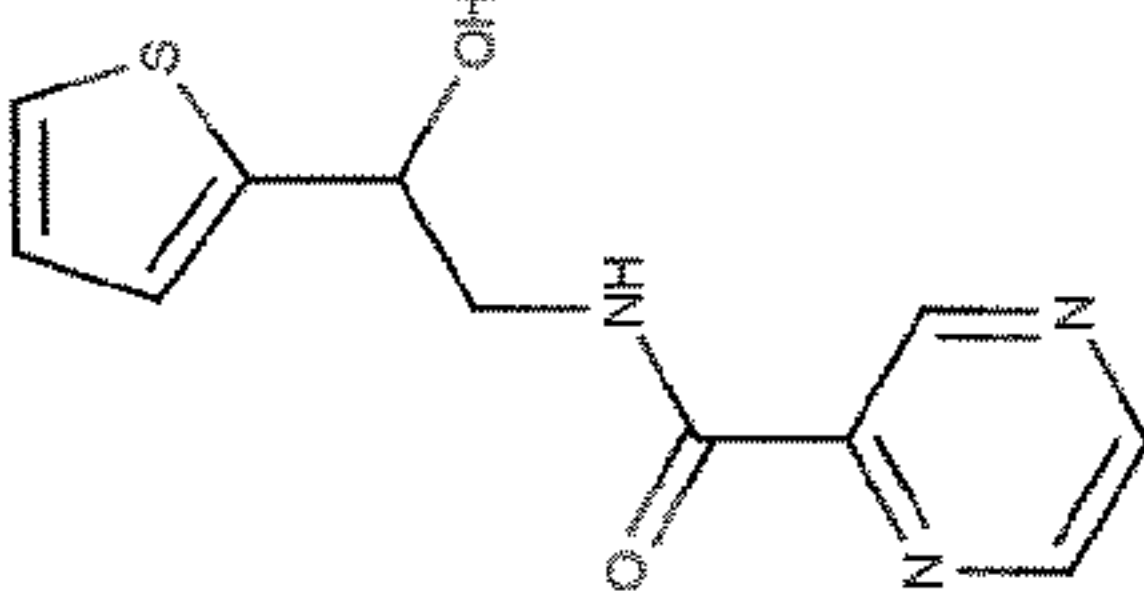
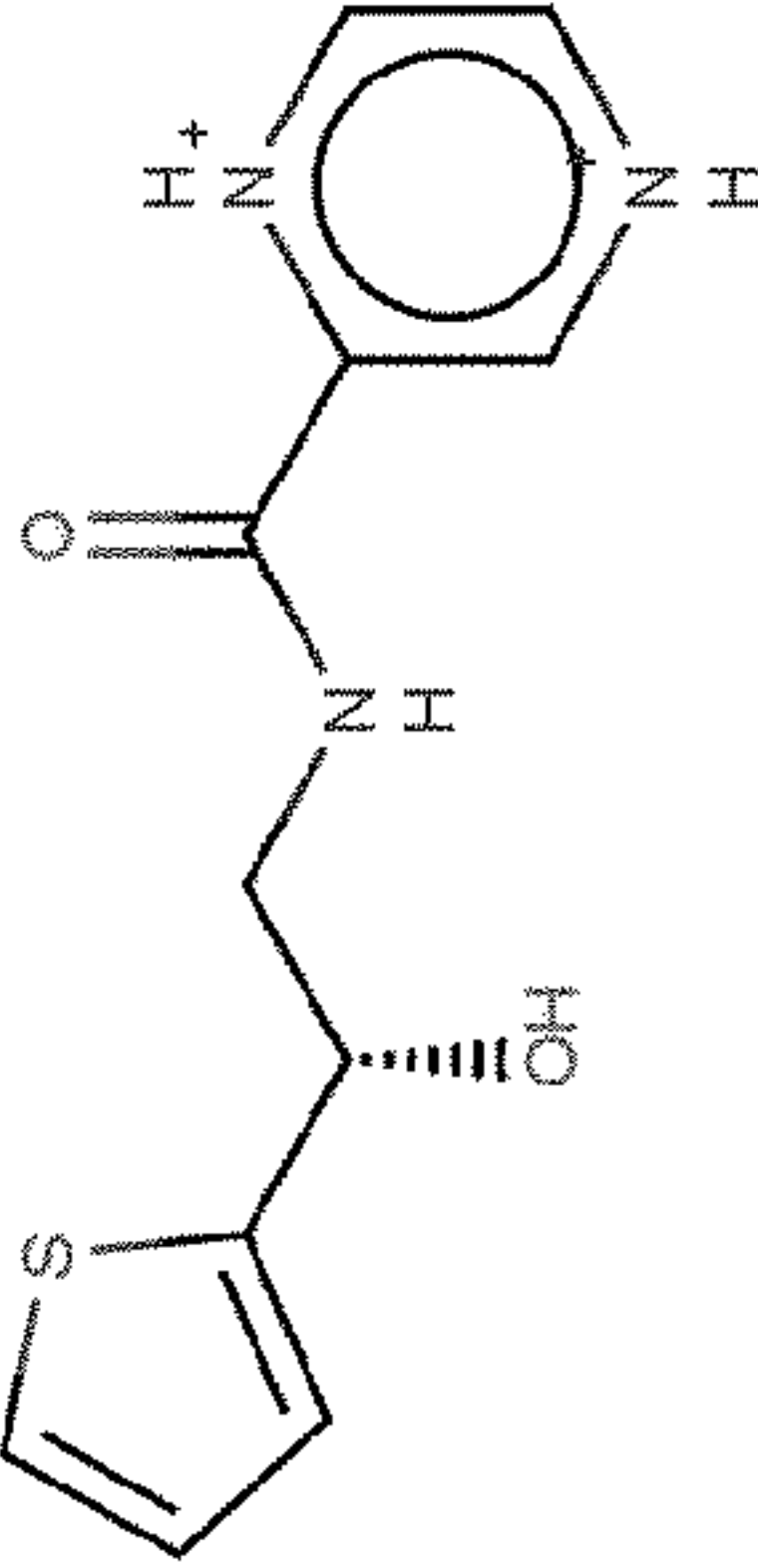
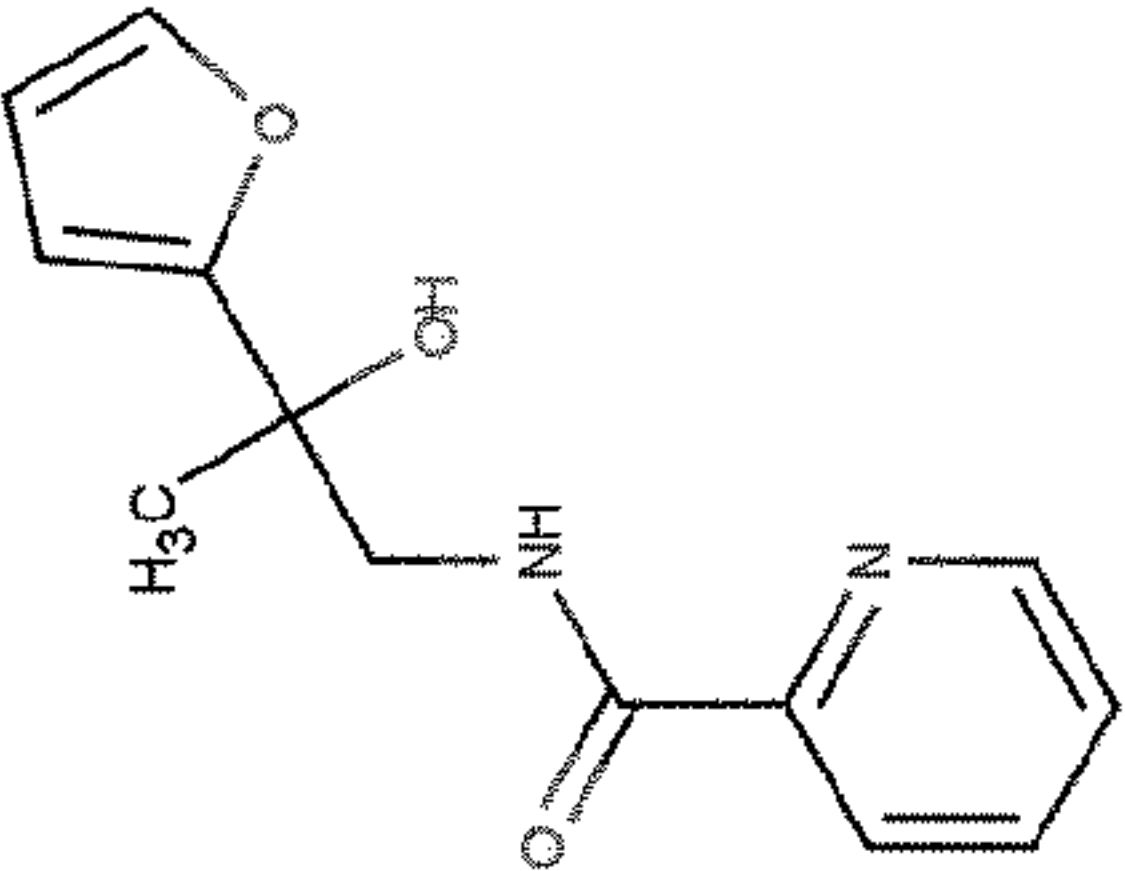
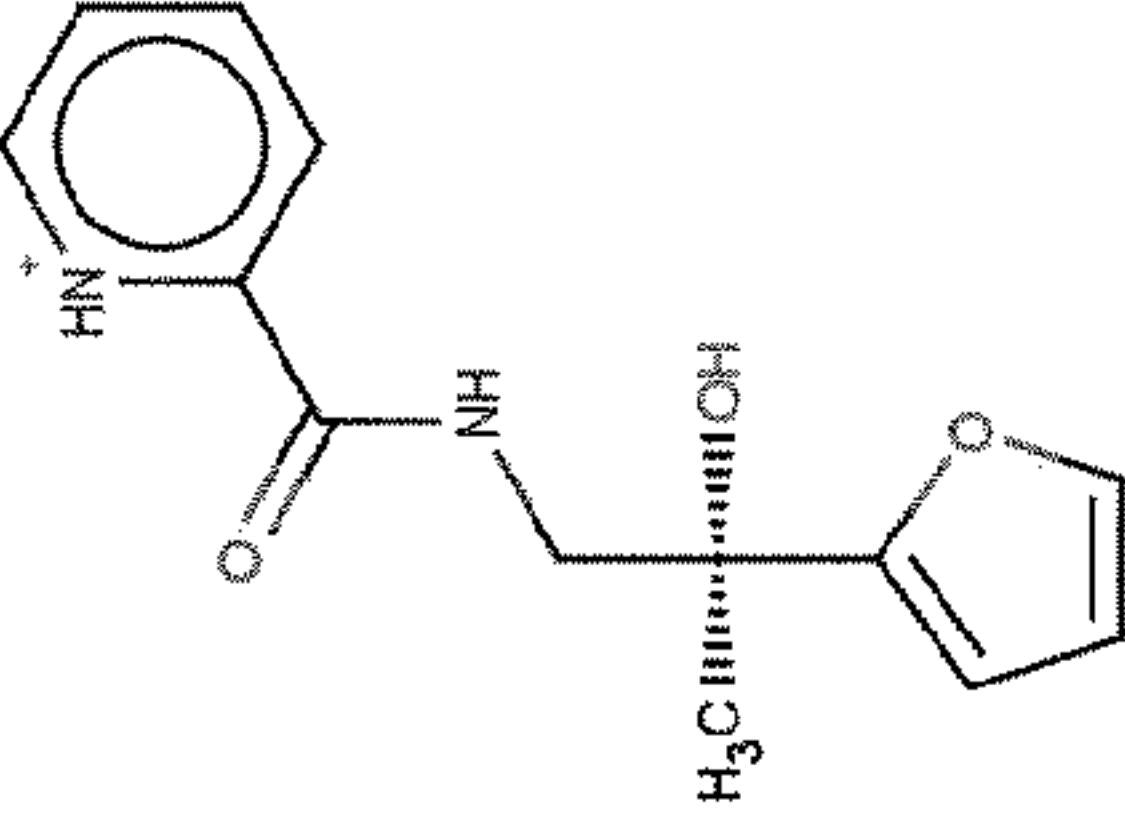
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 229		F5806-0308		F5806-0308_1_1 (1 of 2)
Compound 230		F5828-0220		F5828-0220_3_2 (1 of 14)
Compound 231		F5831-3069		F5831-3069_2_2 (1 of 8)

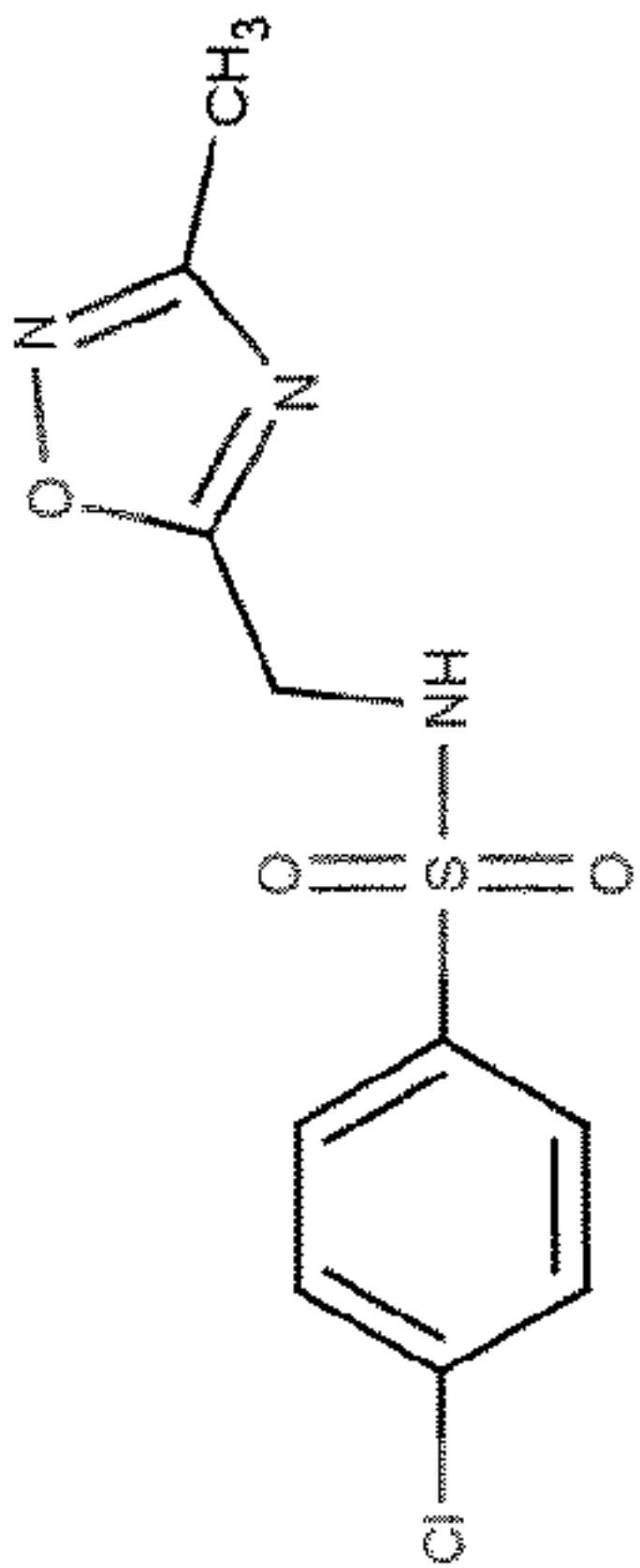
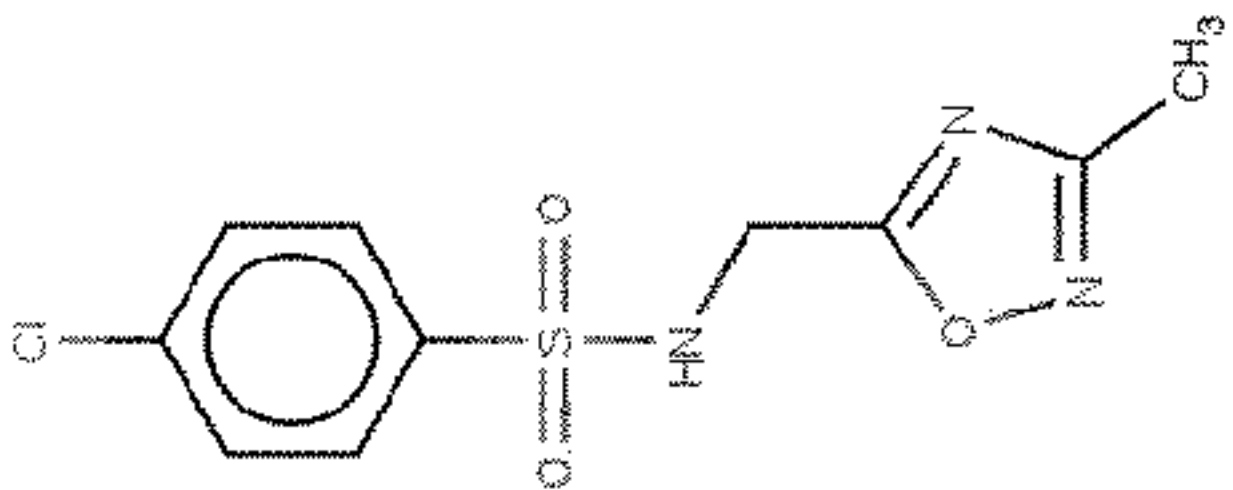
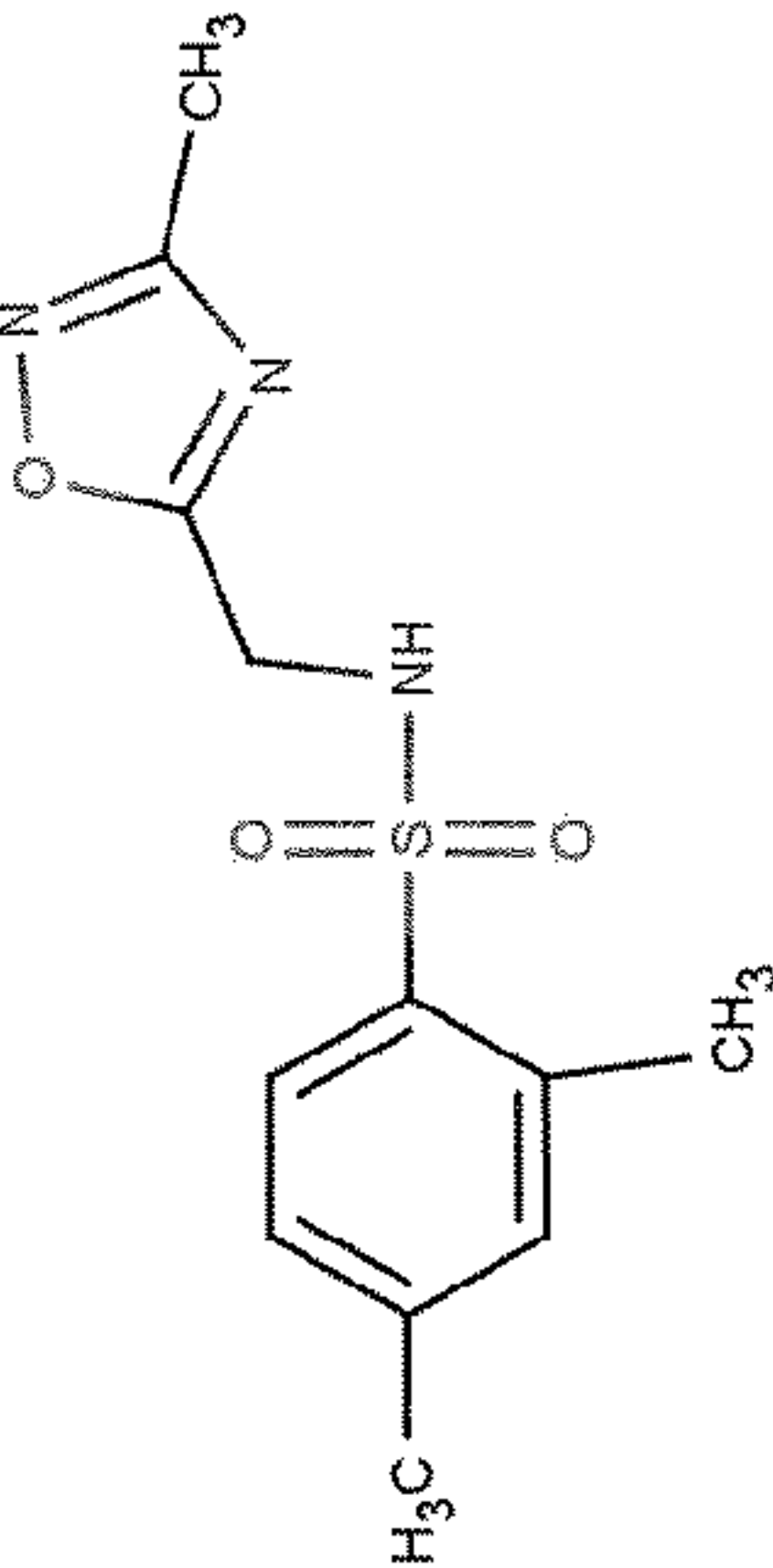
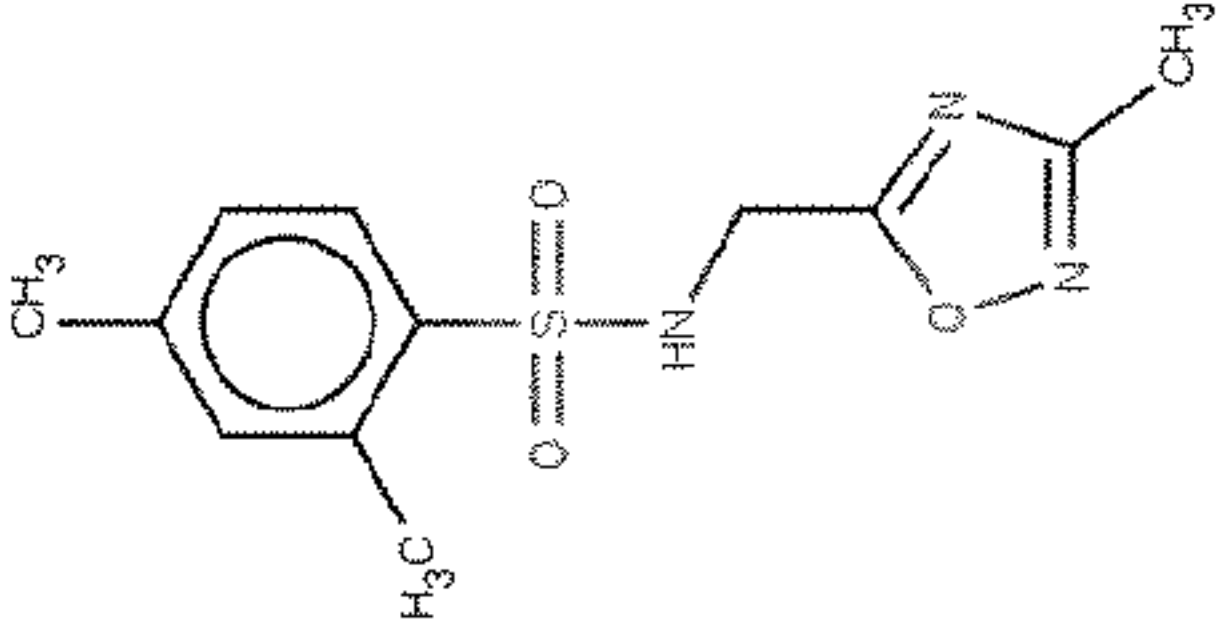
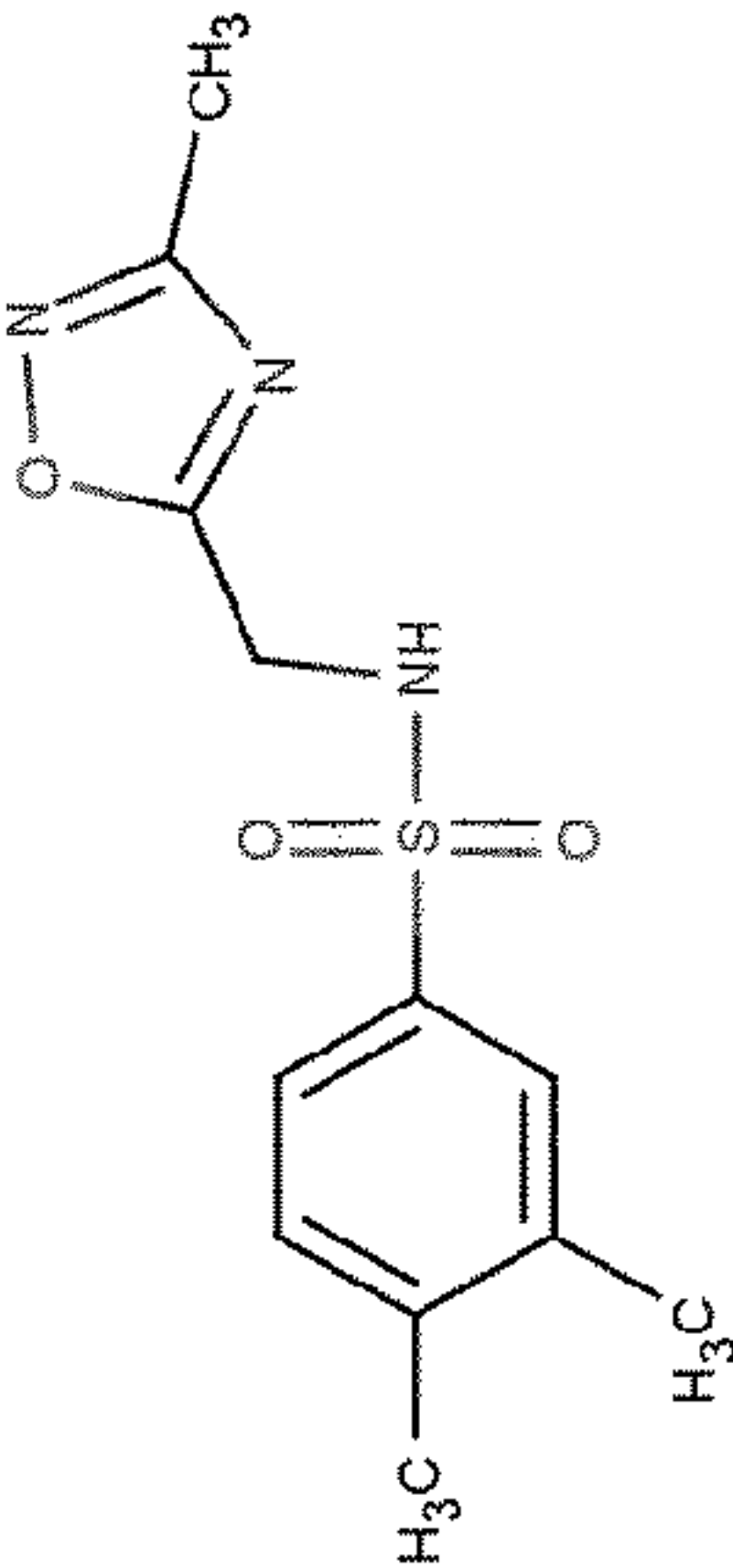
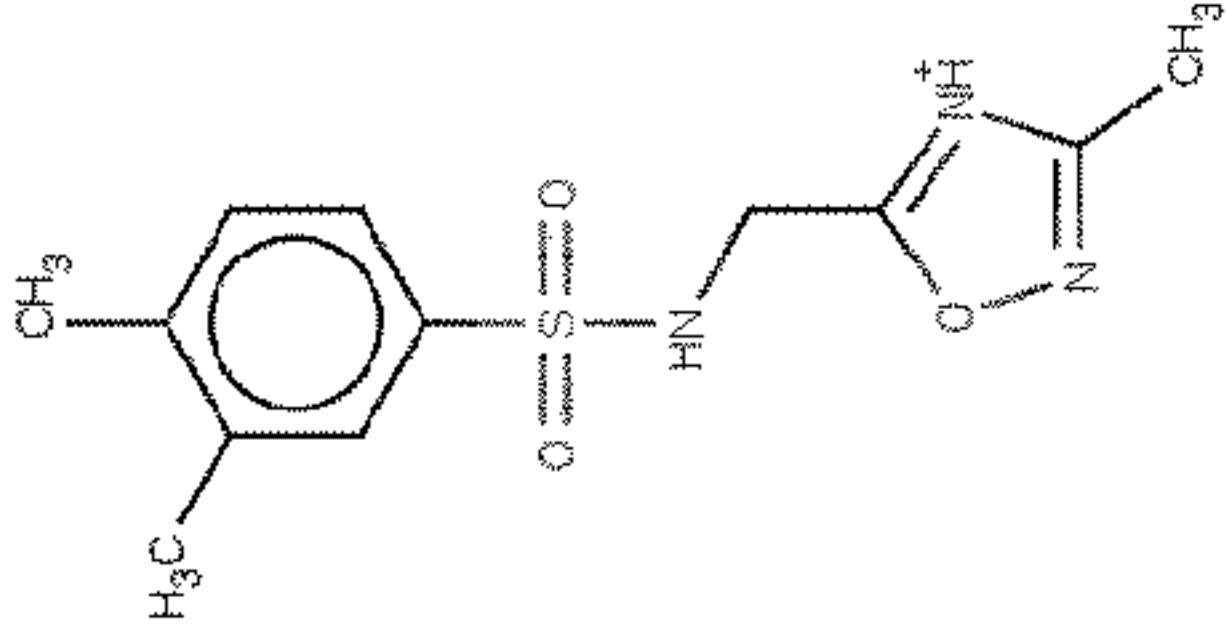
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 232		F5831-6204		F5831-6204_1_1 (1 of 4)
Compound 233		F5856-0133		F5856-0133_3_1 (1 of 2)
Compound 234		F5857-0026		F5857-0026_1 (1 of 2)

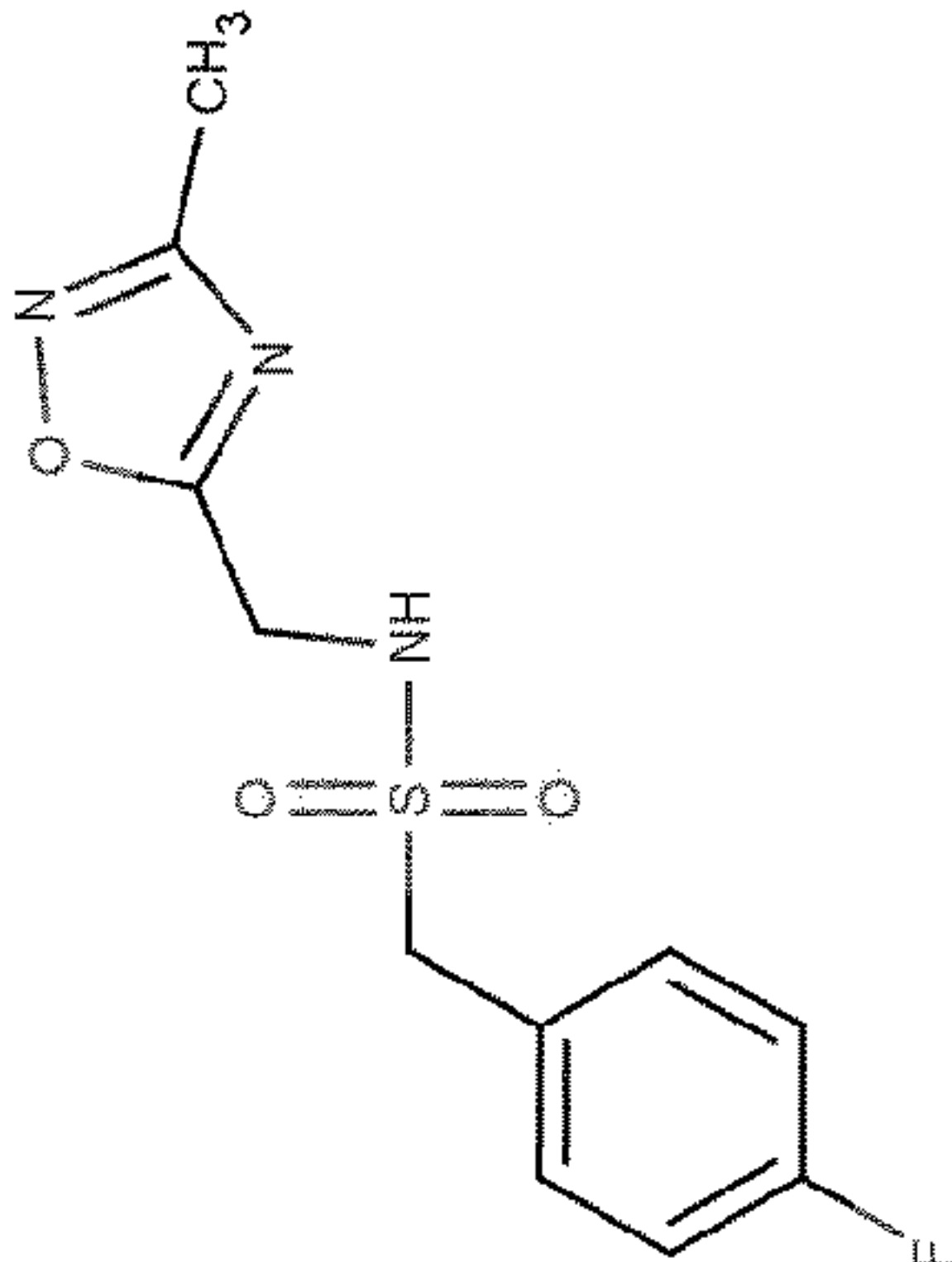
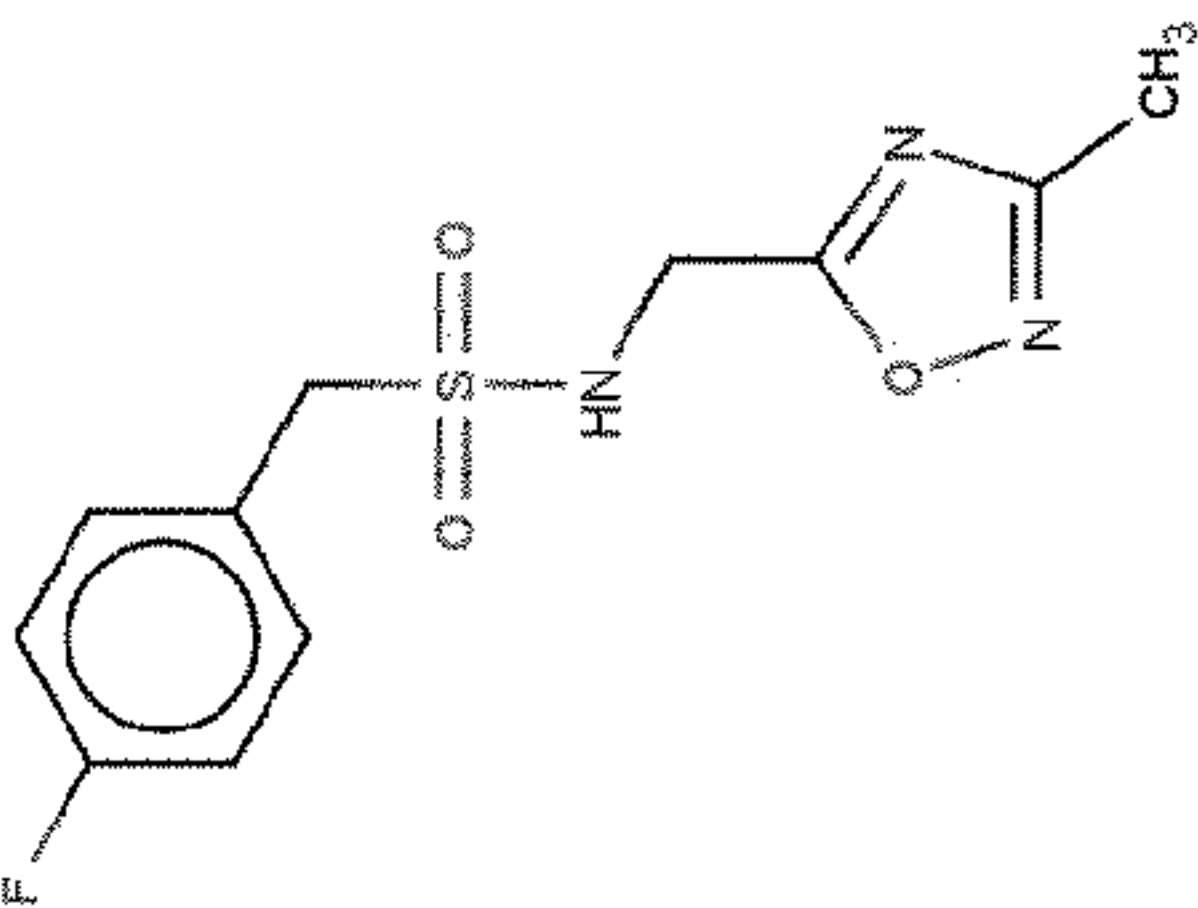
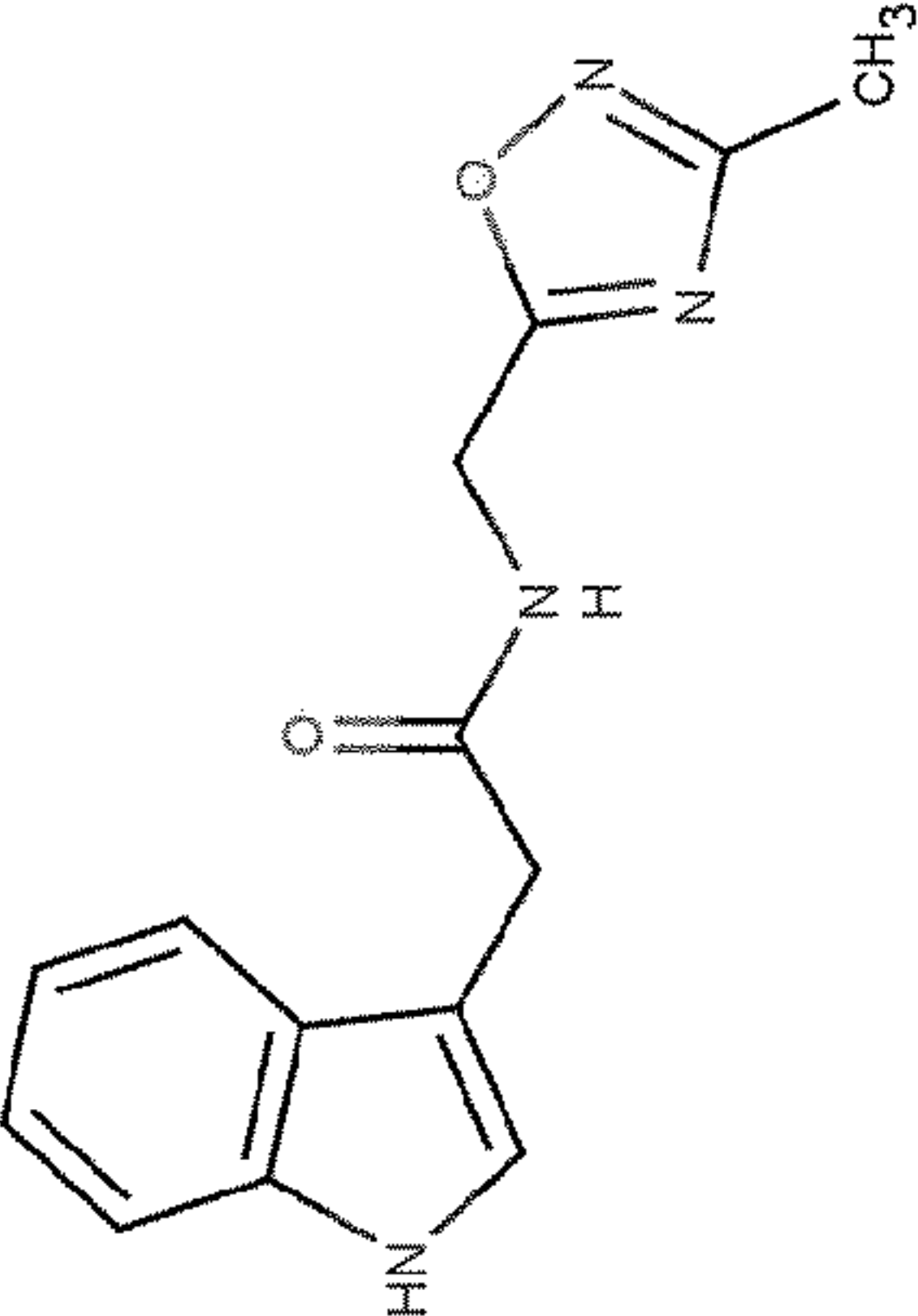
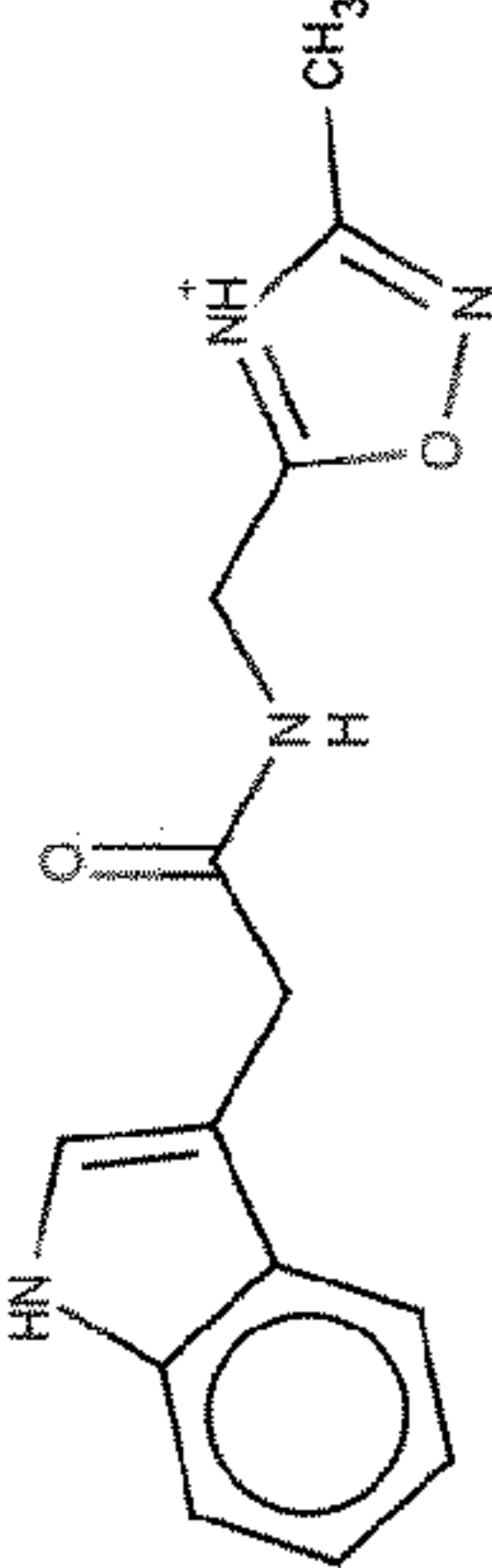
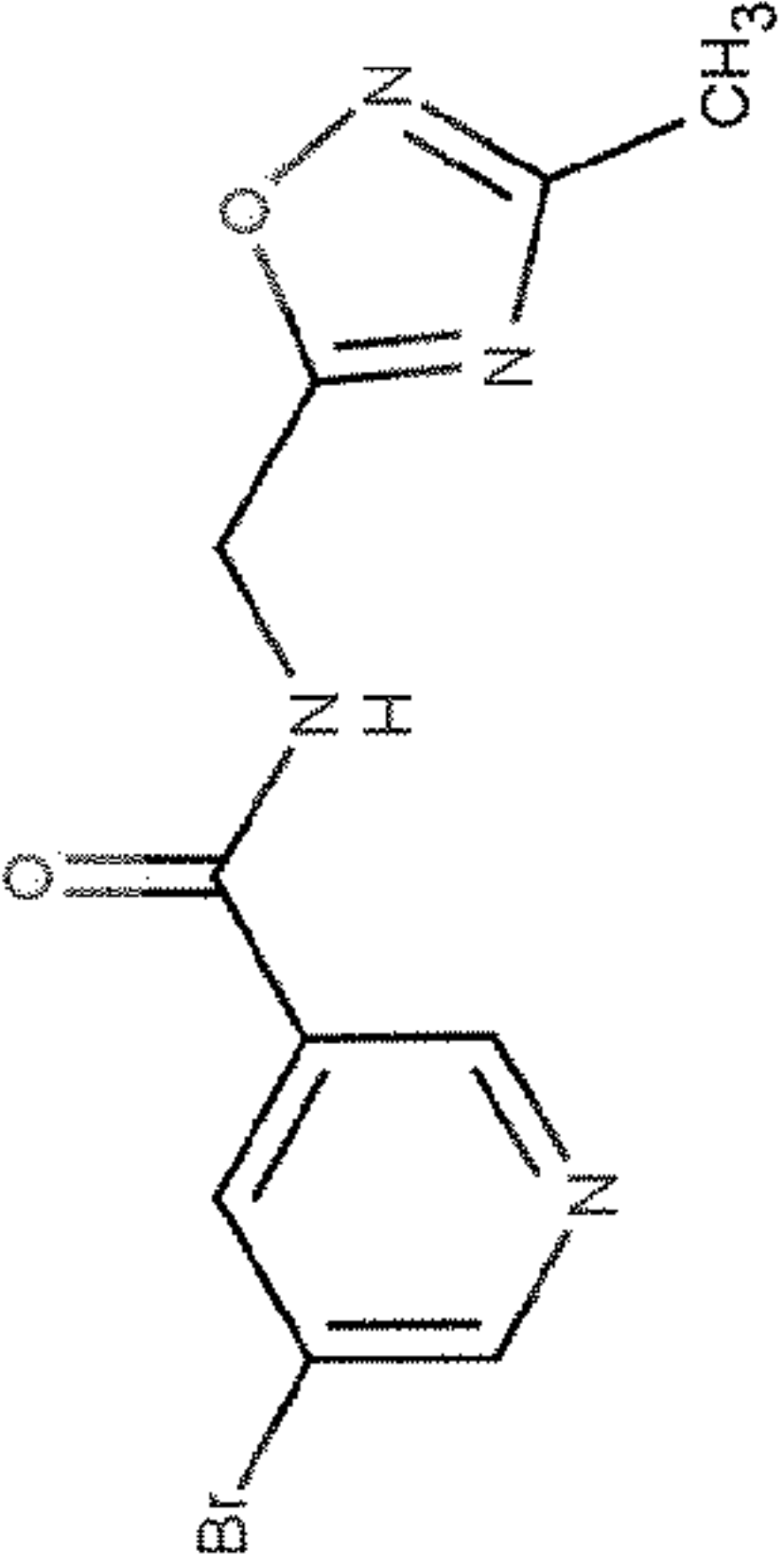
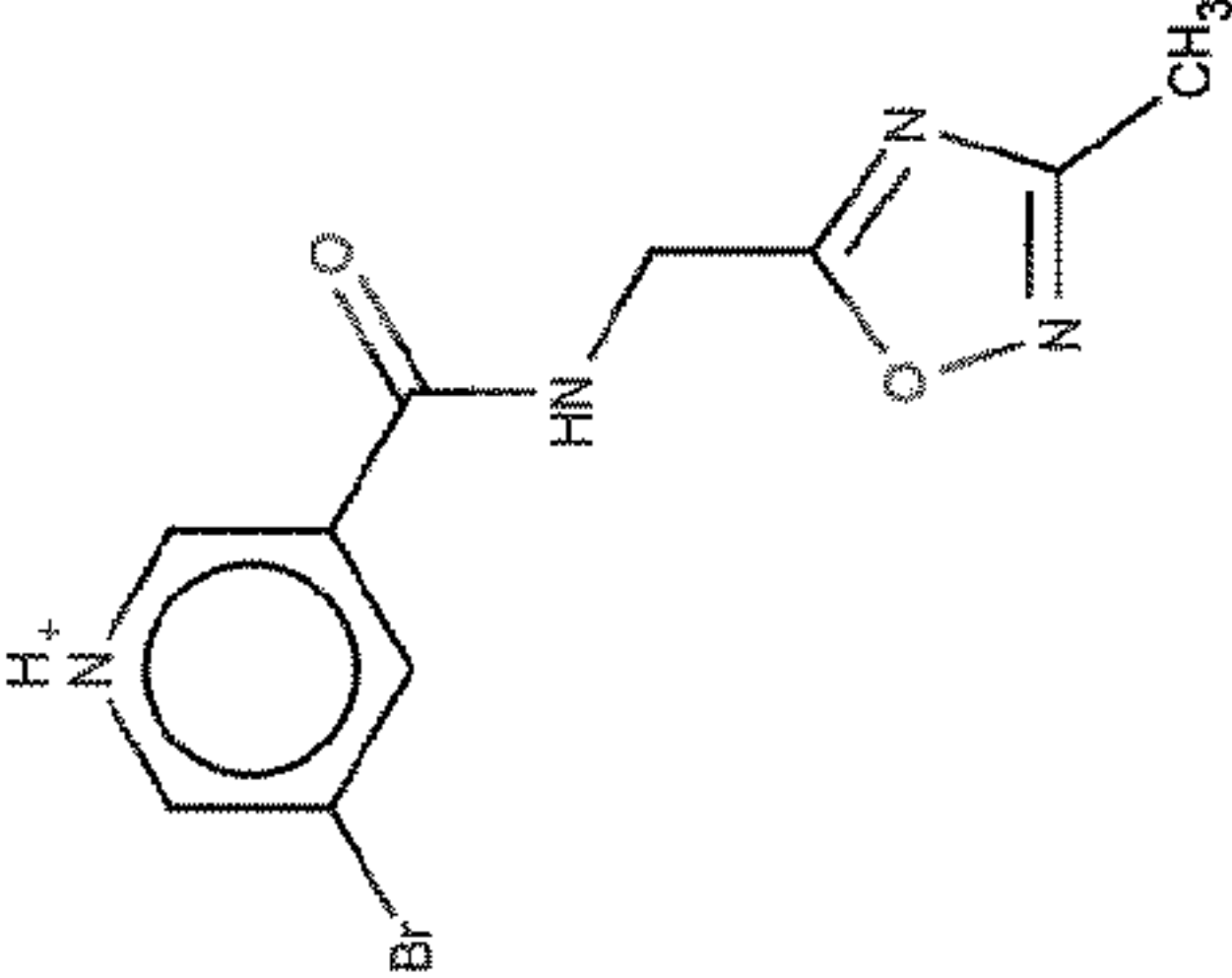
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Compound 235		F5857-0927		F5857-0927_1 (1 of 2)
Compound 236		F5857-1145		F5857-1145_1_1 (1 of 8)
Compound 237		F5857-1524		F5857-1524_1_1 (1 of 8)

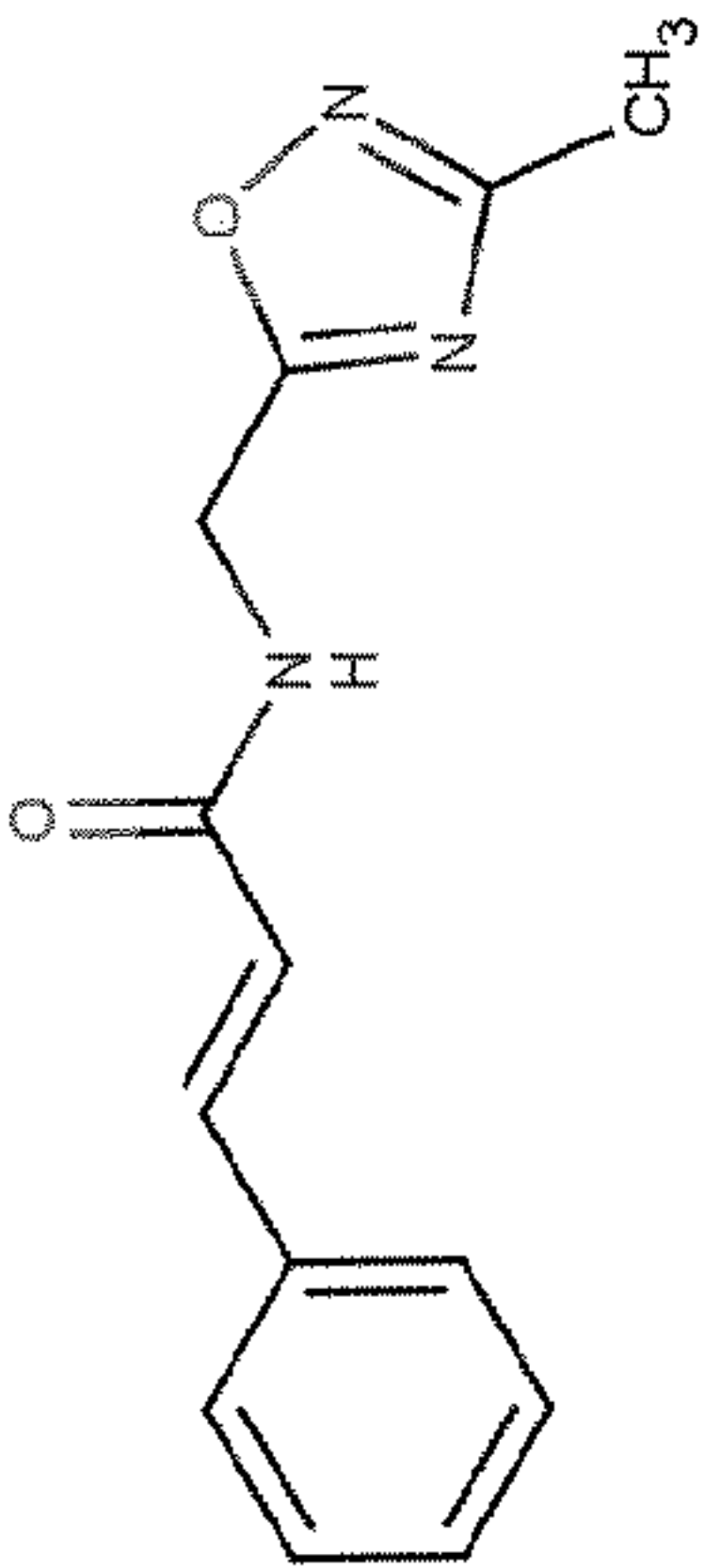
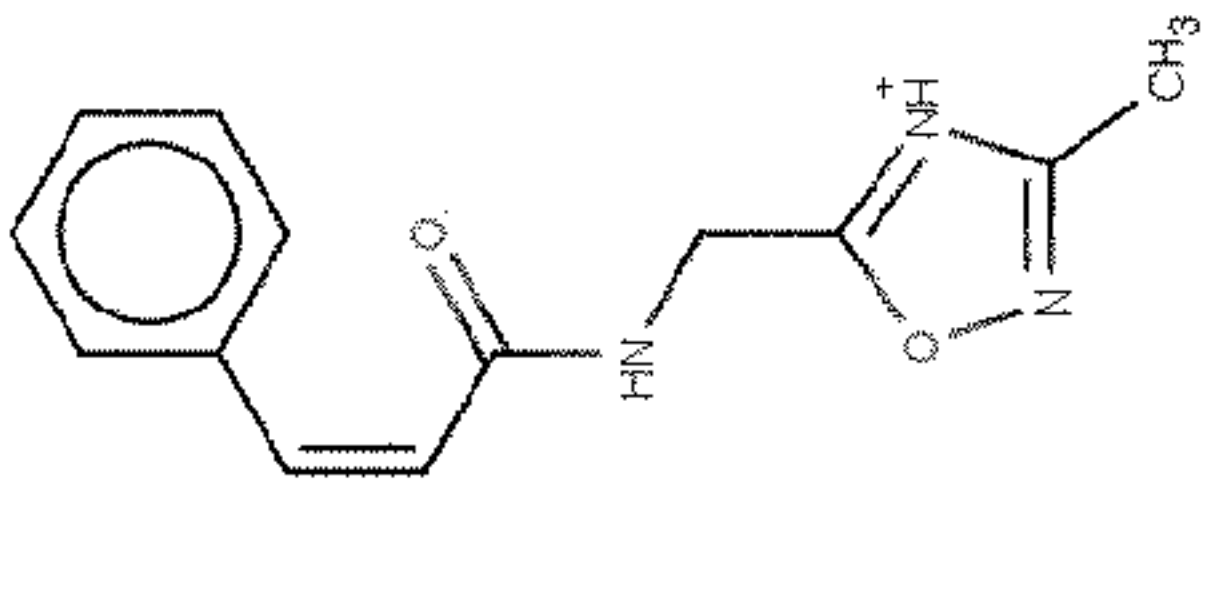
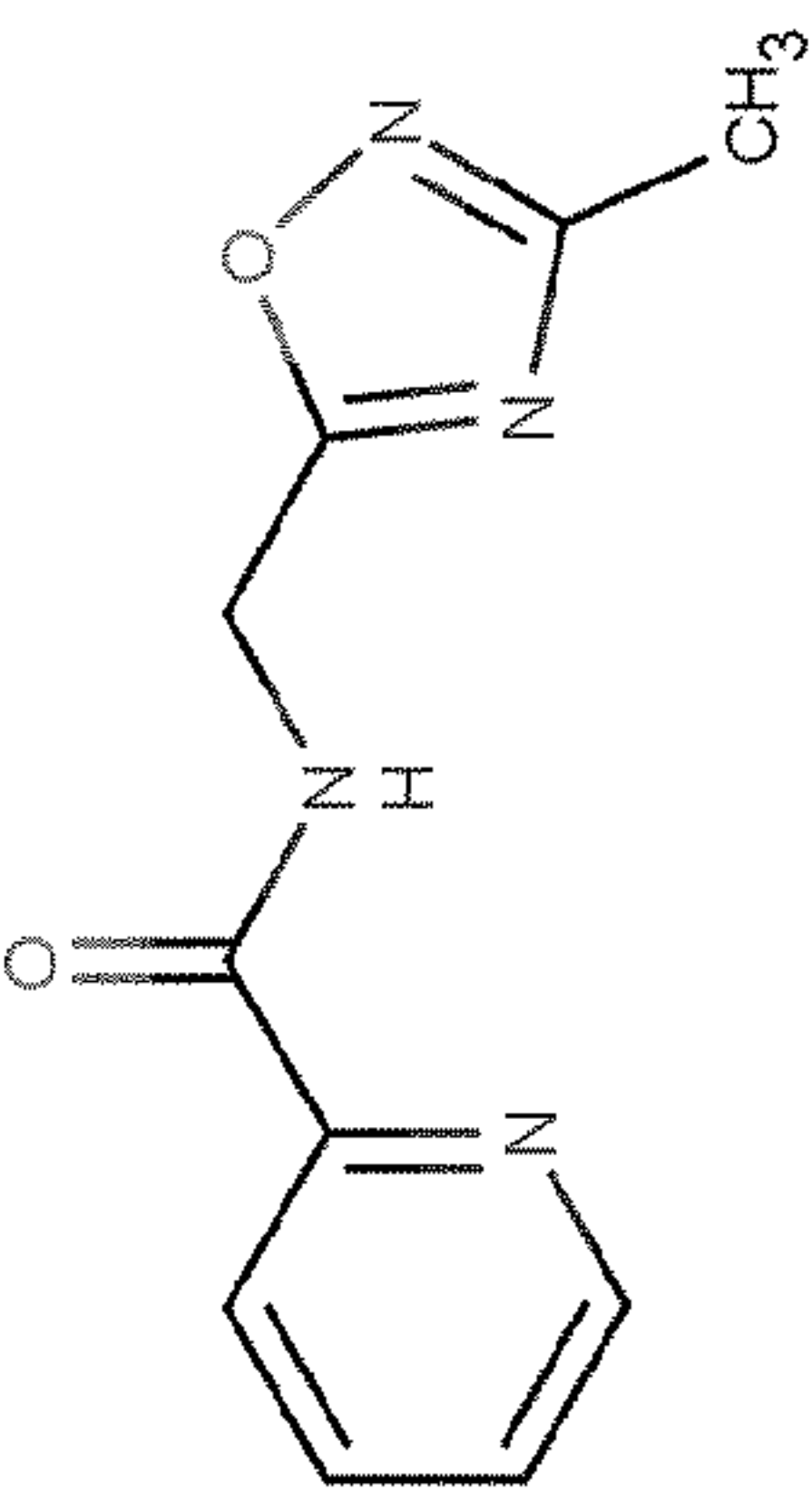
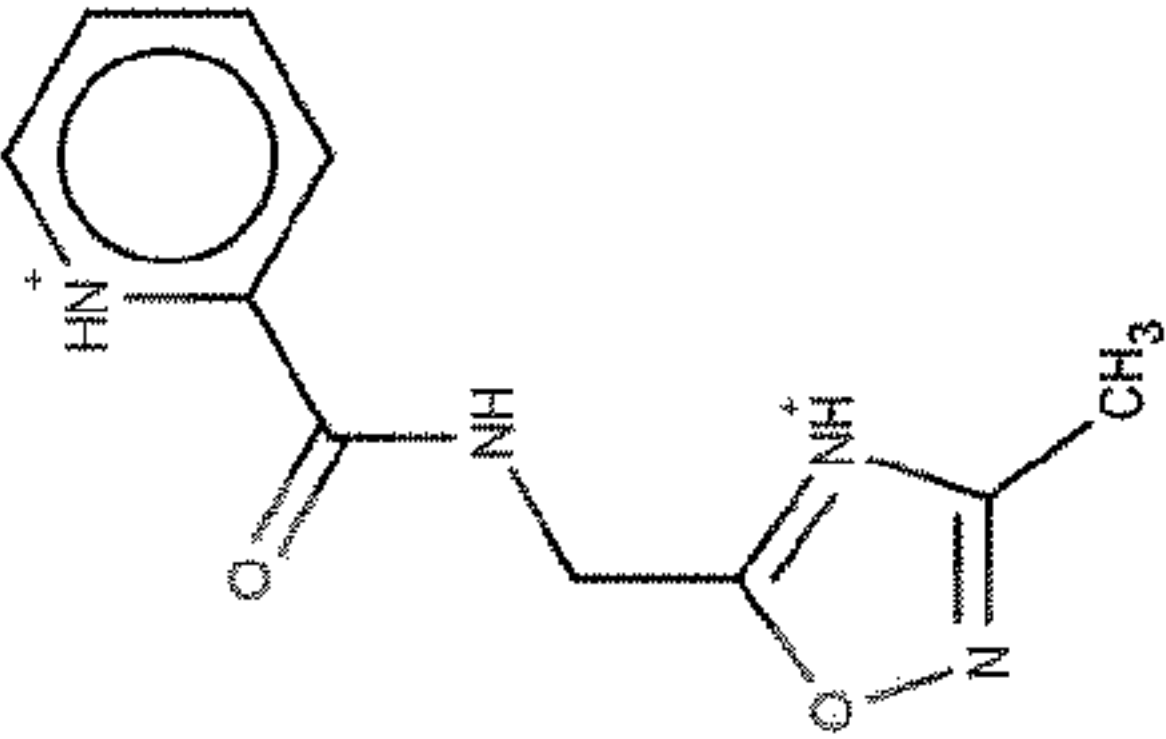
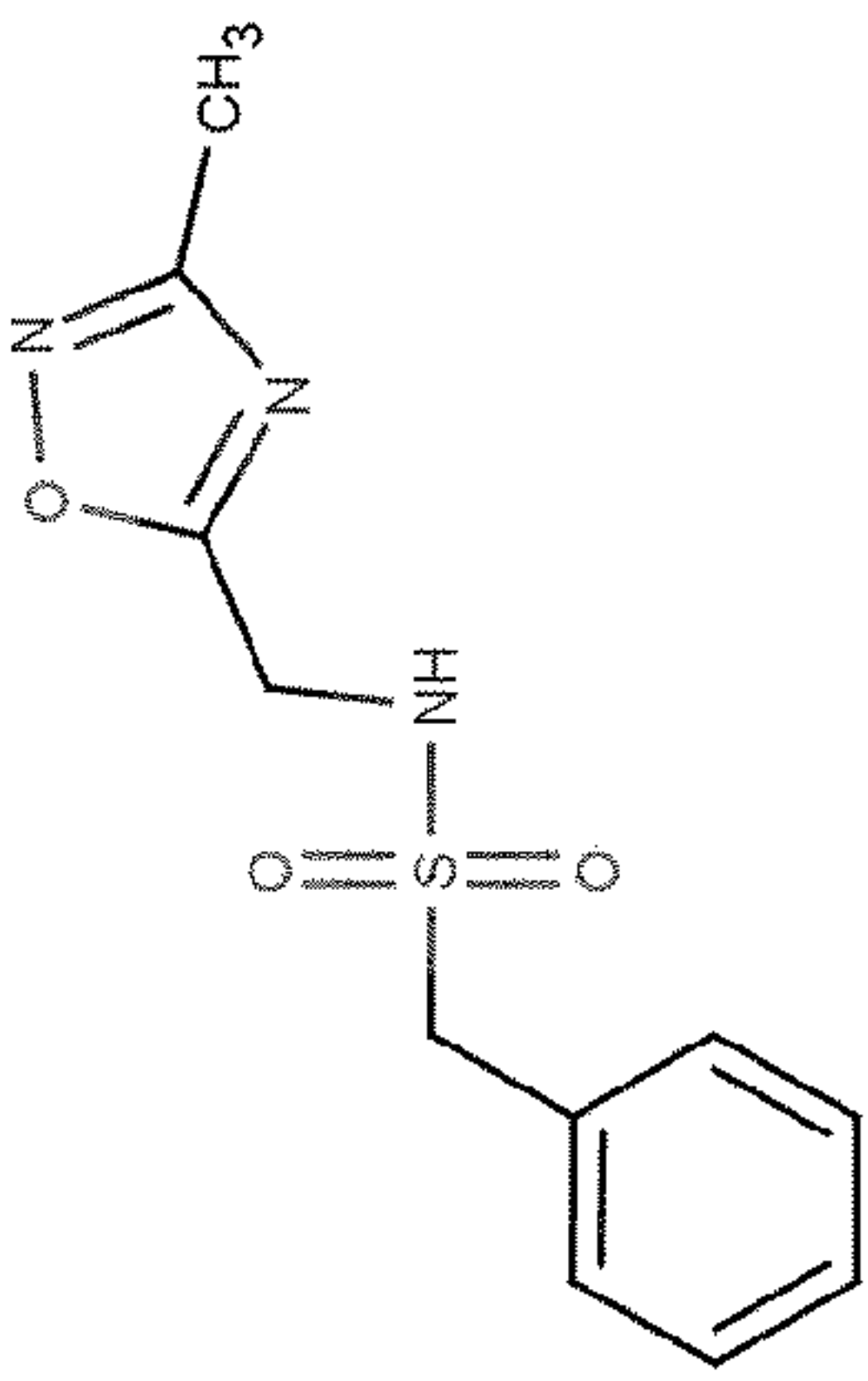
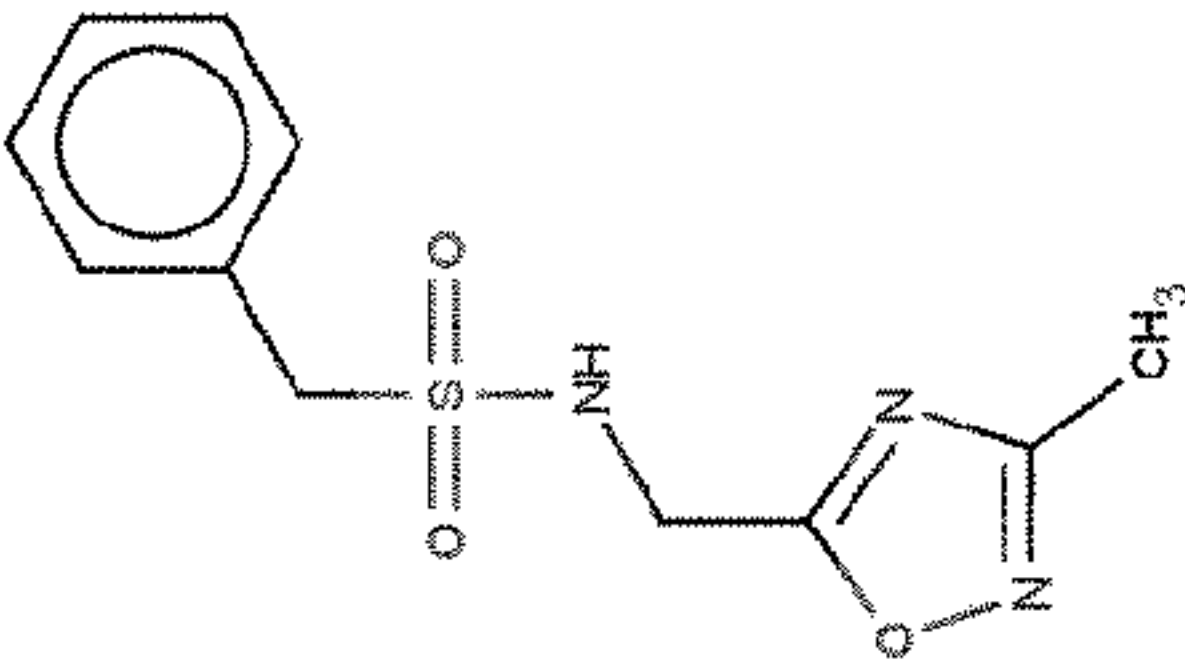
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 238		F5857-4935		F5857-4935_1_2 (1 of 8)
Compound 239		F5857-5314		F5857-5314_1_2 (1 of 8)
Compound 240		F5857-5693		F5857-5693_1_1 (1 of 8)

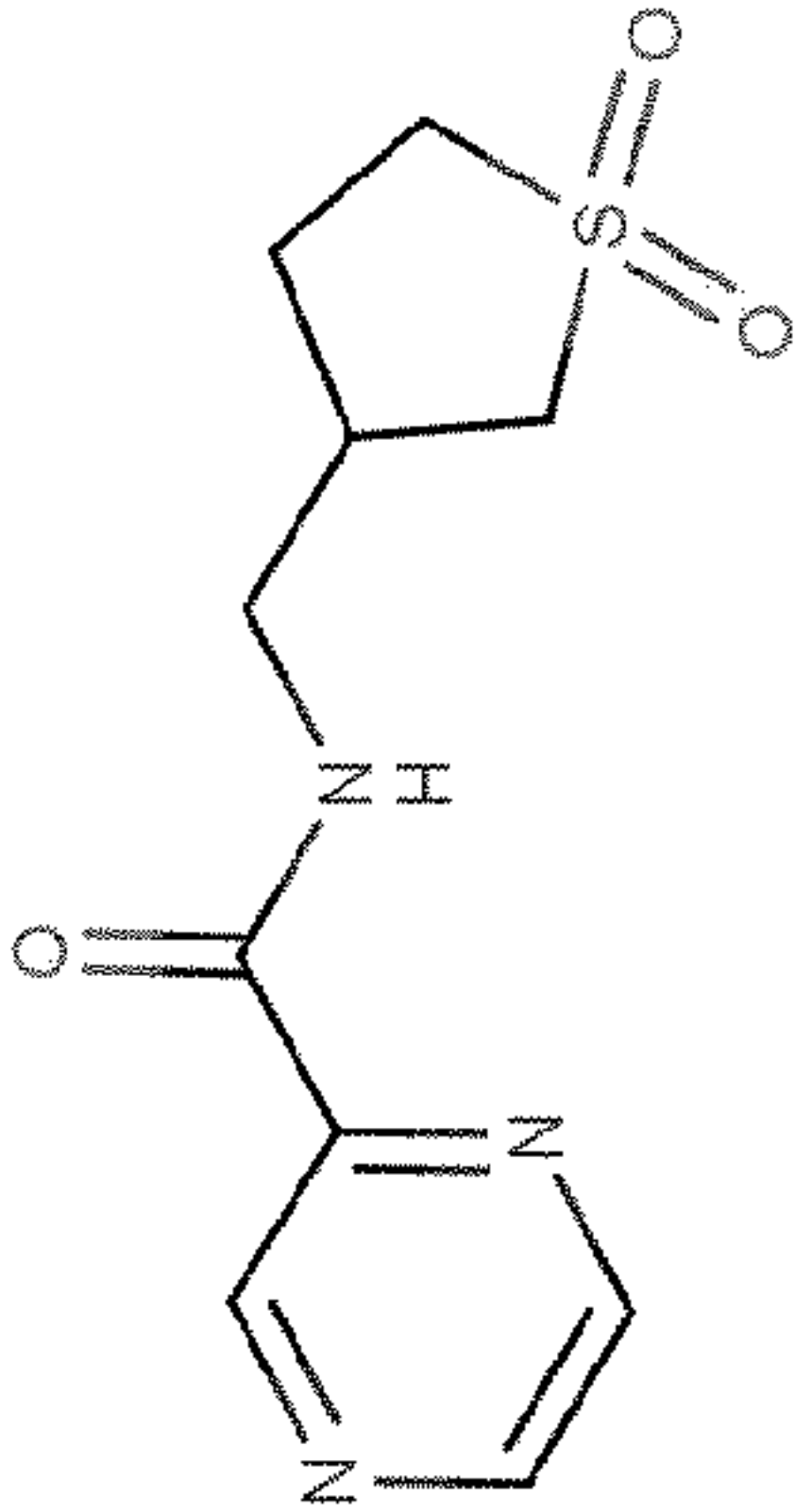
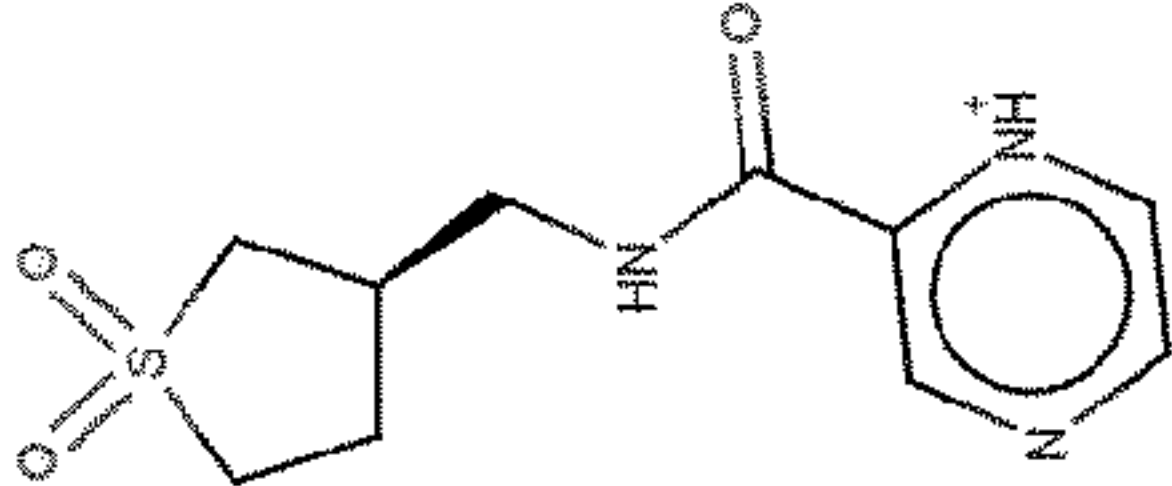
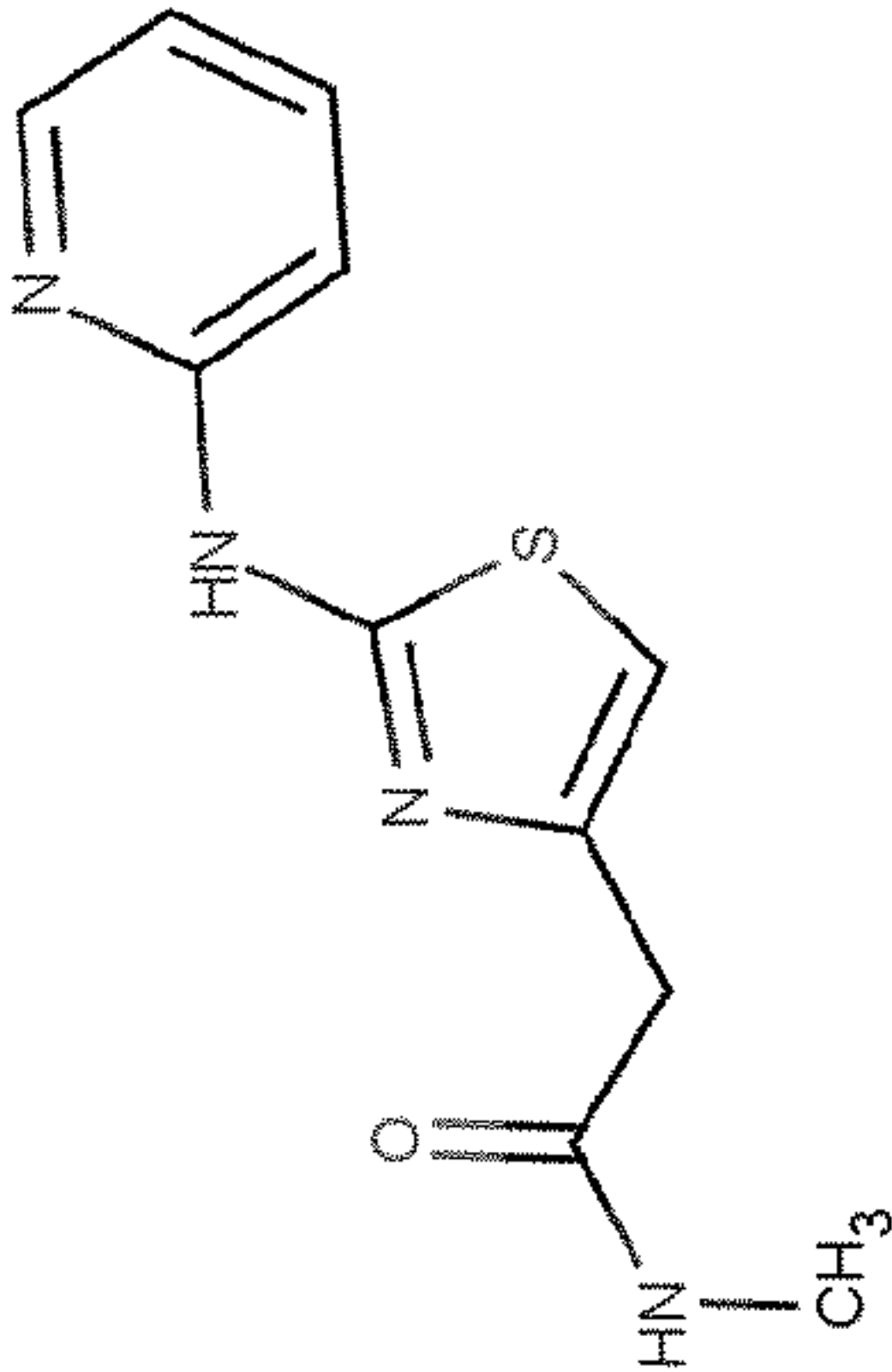
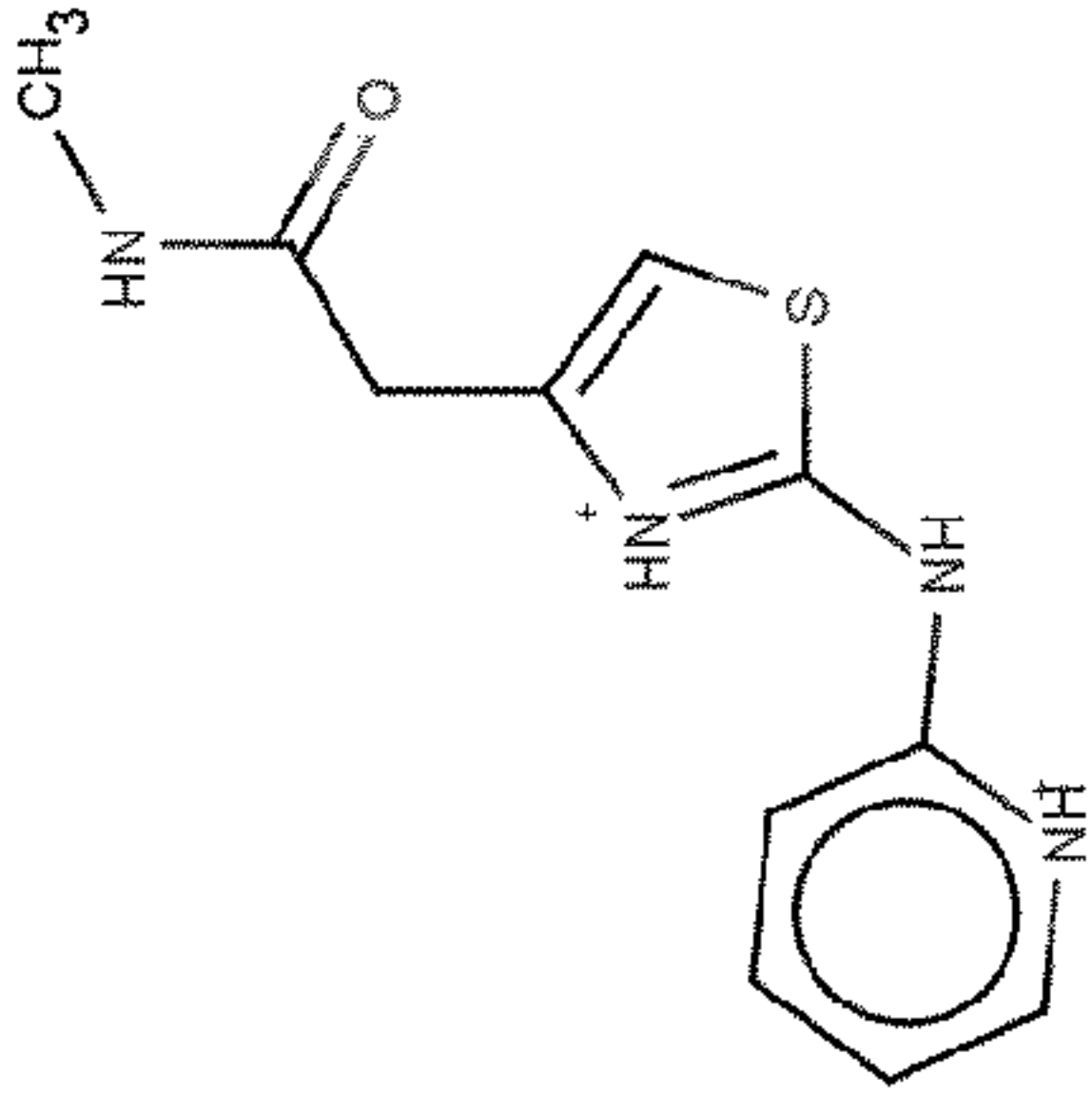
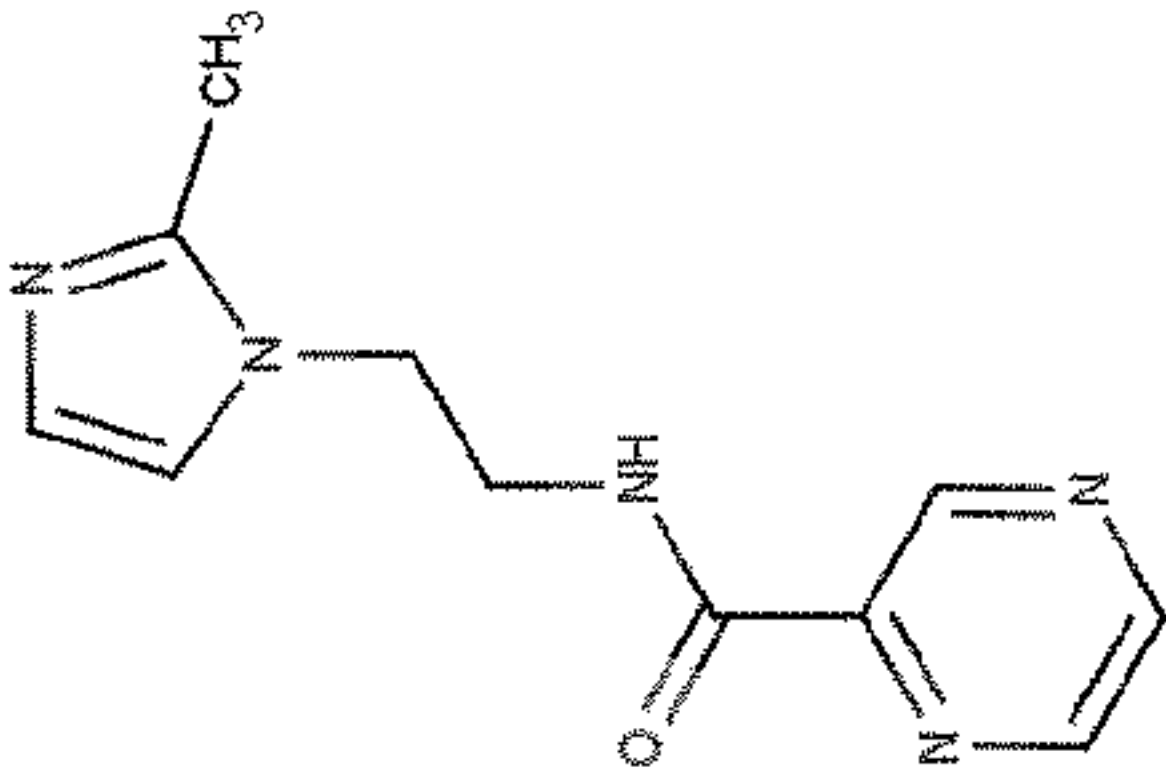
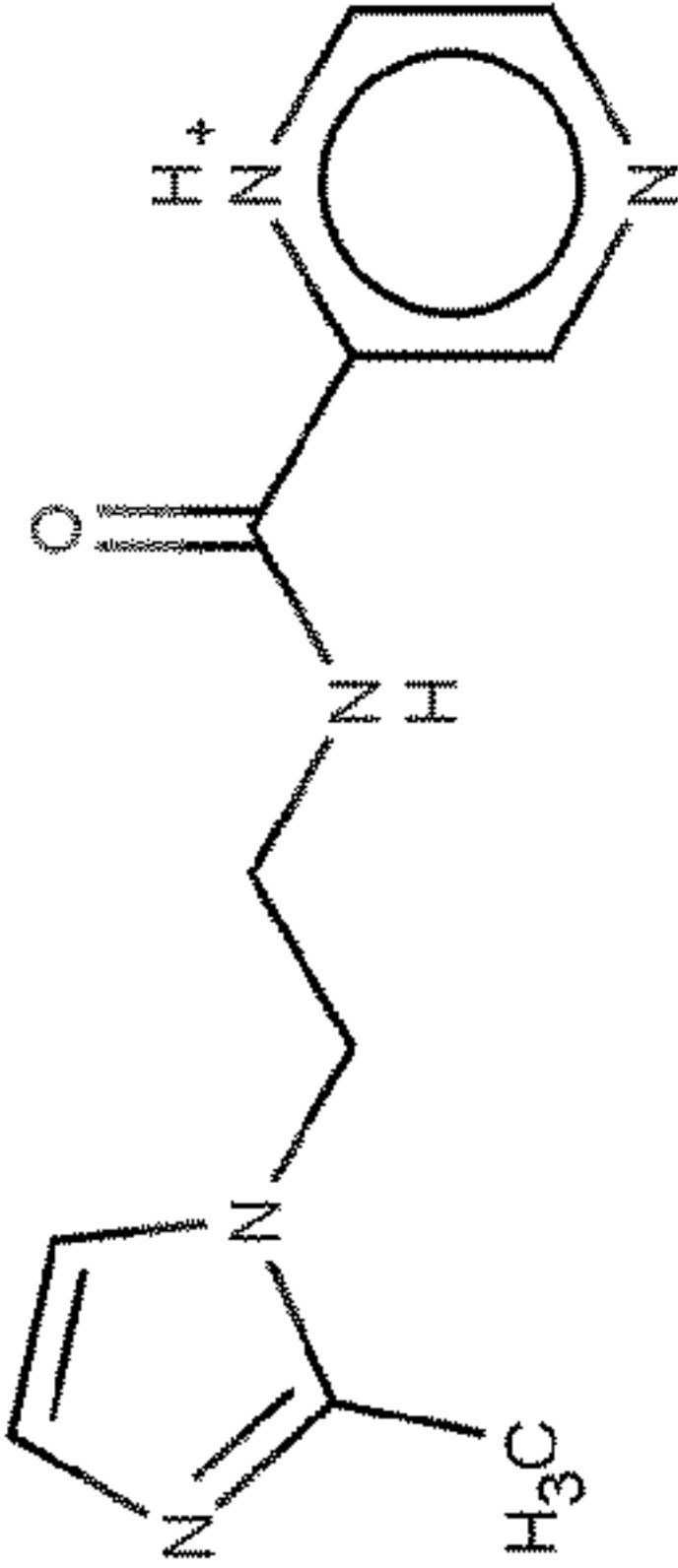
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 241		F5857-5862		F5857-5862_1 (1 of 2)
Compound 242		F5857-6072		F5857-6072_1_1 (1 of 4)
Compound 243		F5857-6266		F5857-6266_1 (1 of 1)

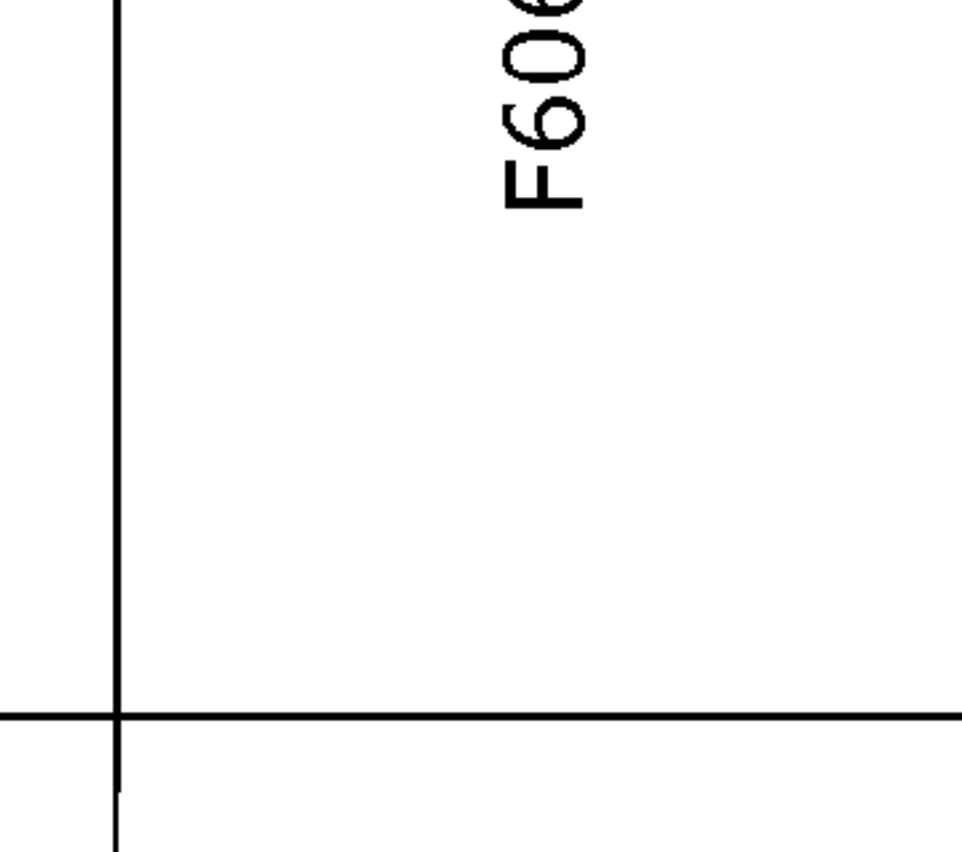
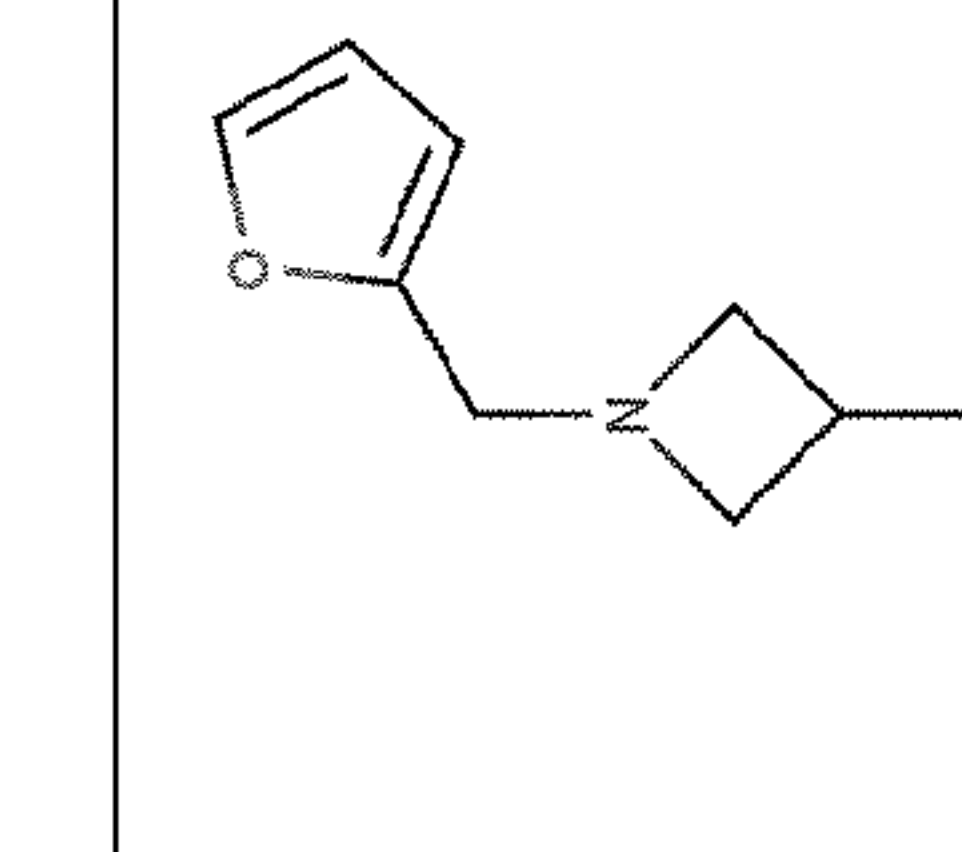
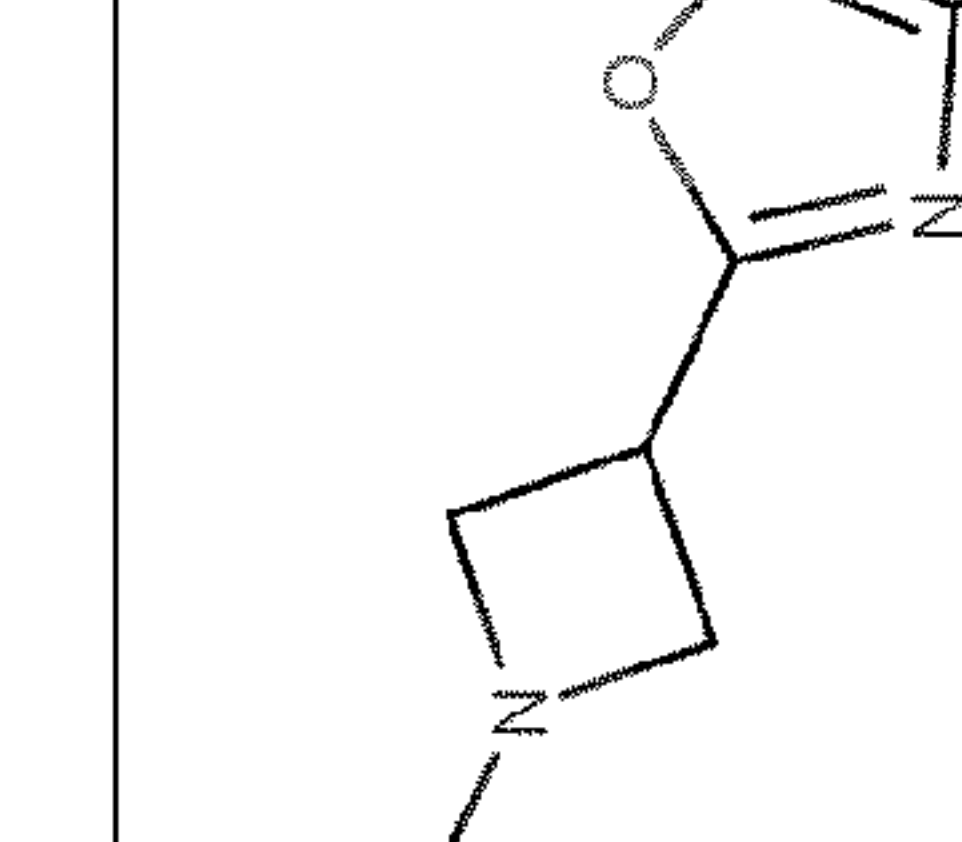
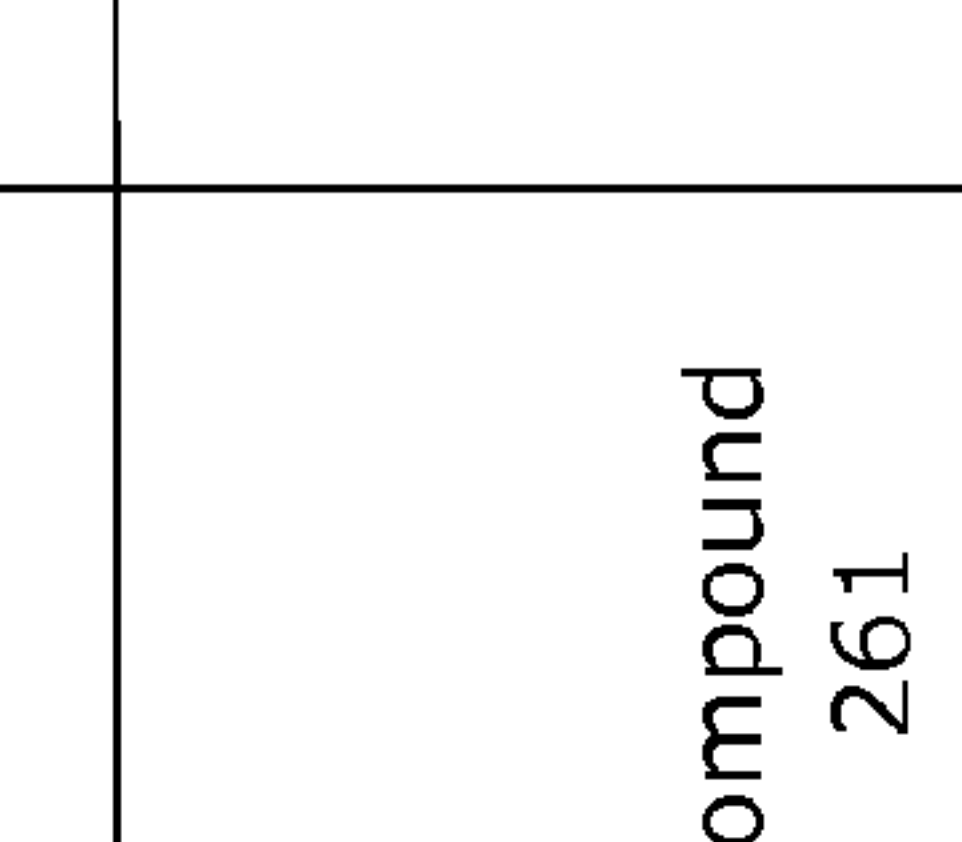
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 244		F5857-7563		F5857-7563_1_2 (1 of 4)
Compound 245		F5857-7901		F5857-7901_3_2 (1 of 8)
Compound 246		F5857-9048		F5857-9048_1_2 (1 of 4)

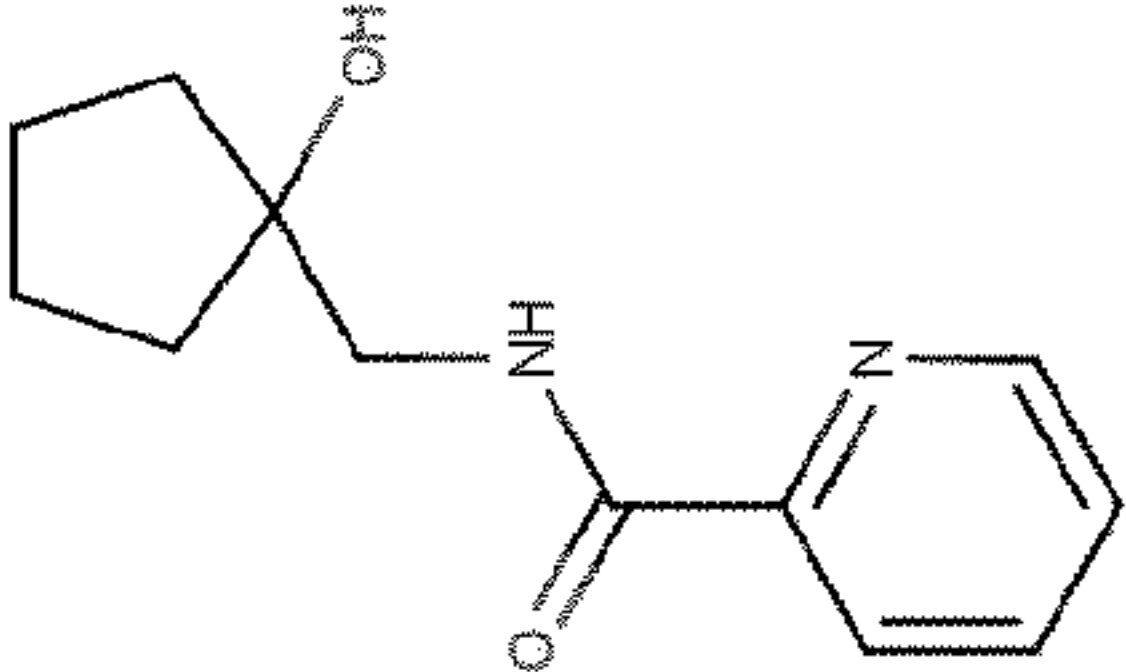
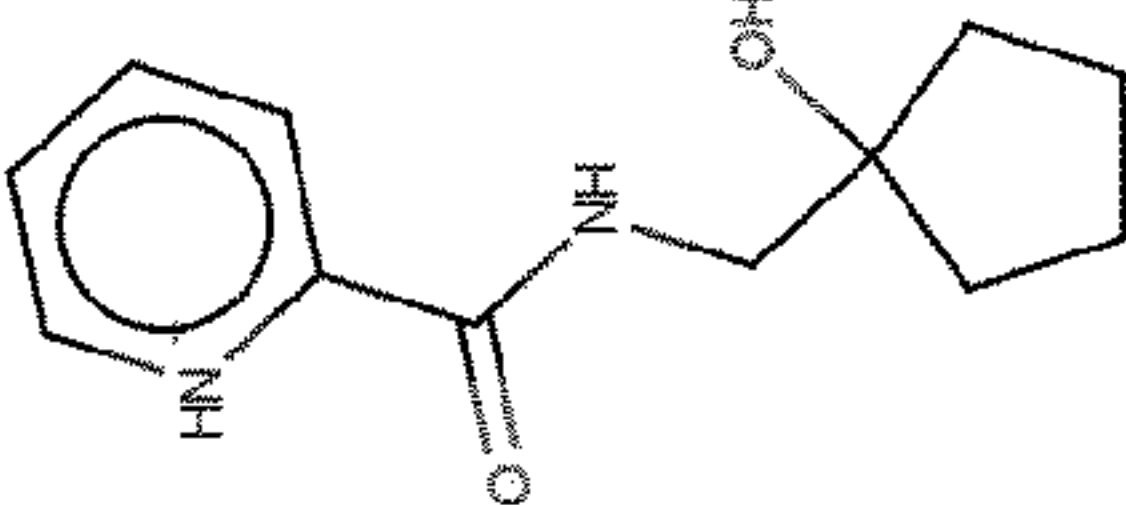
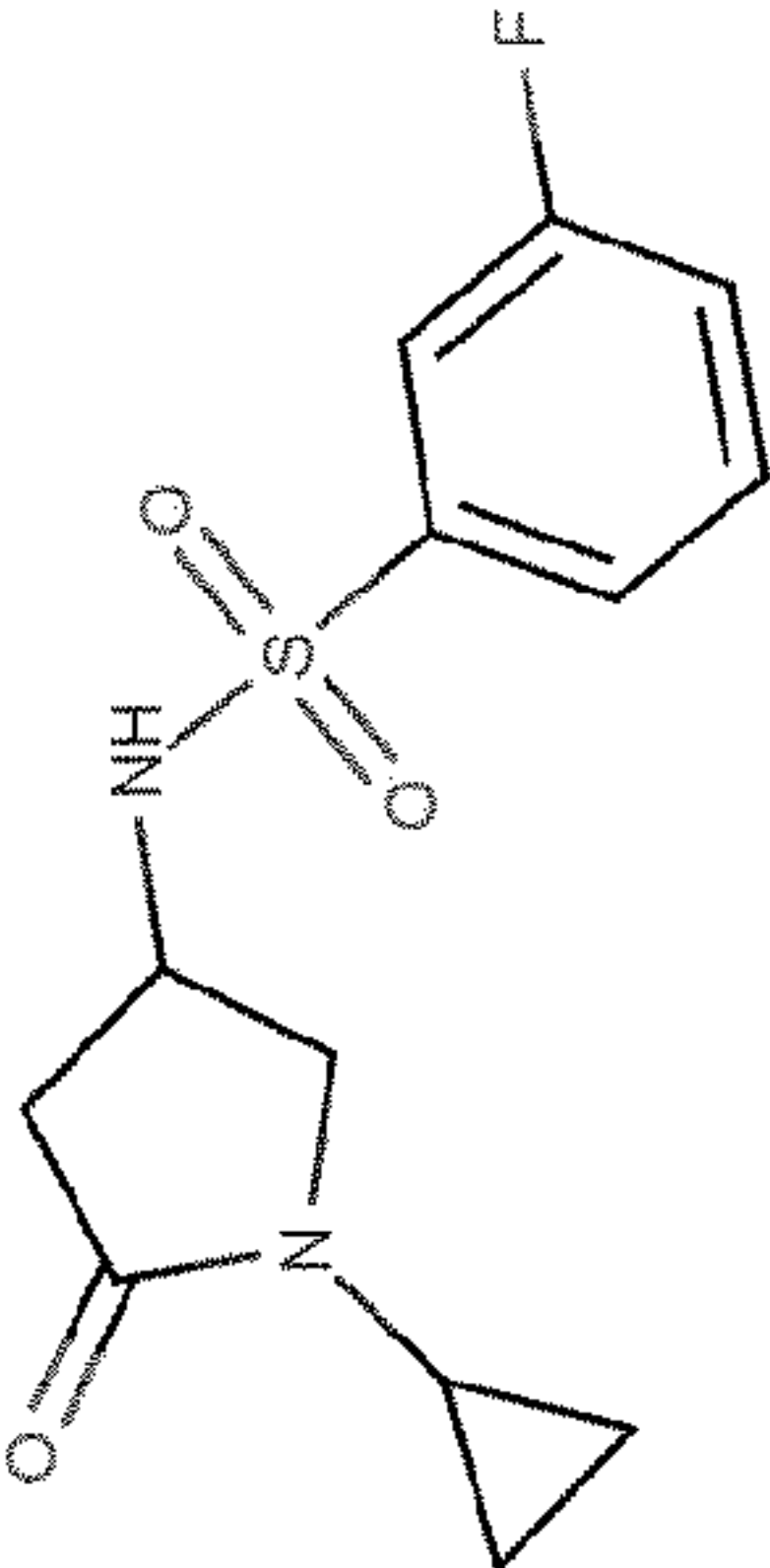
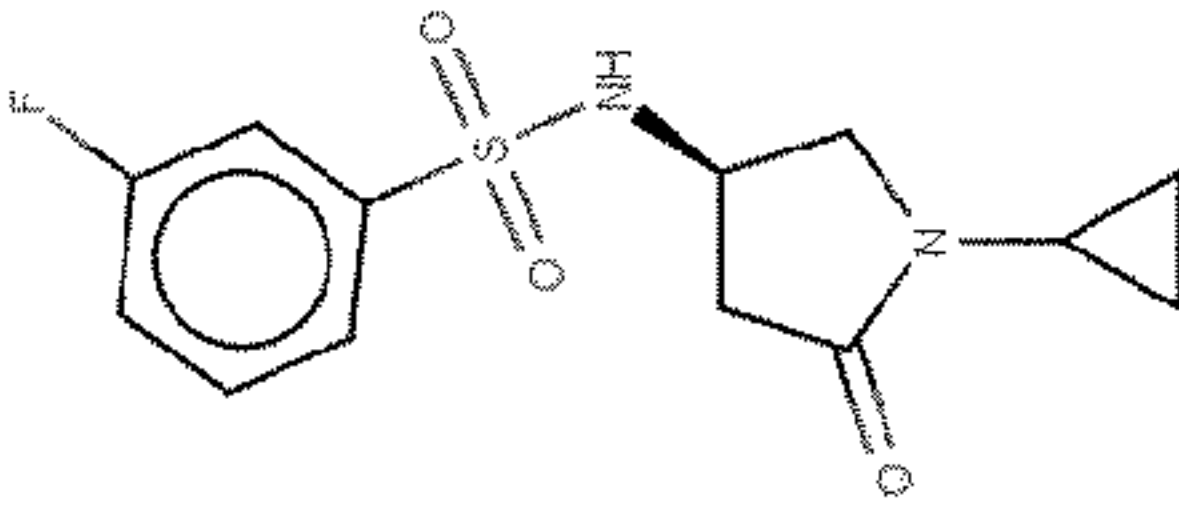
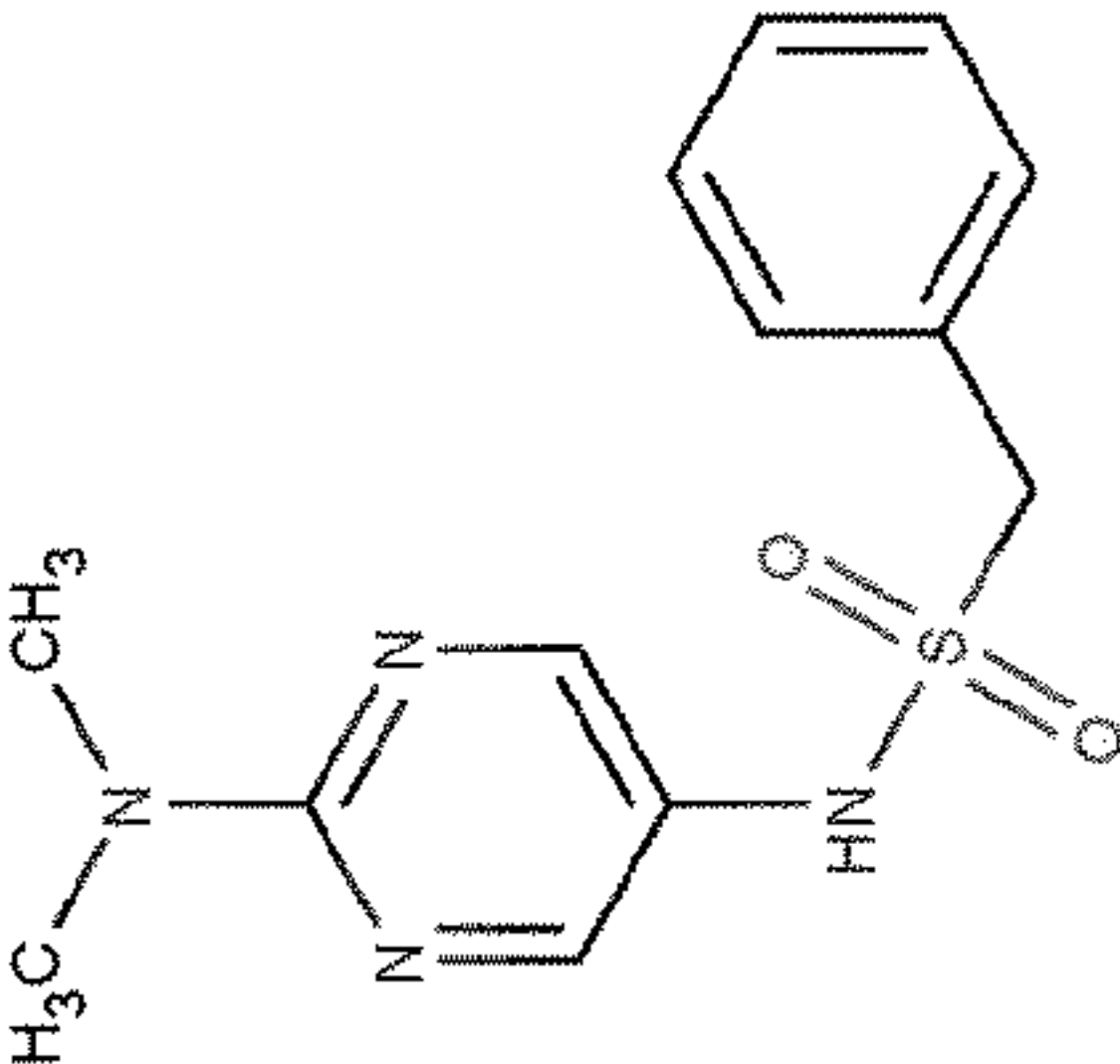
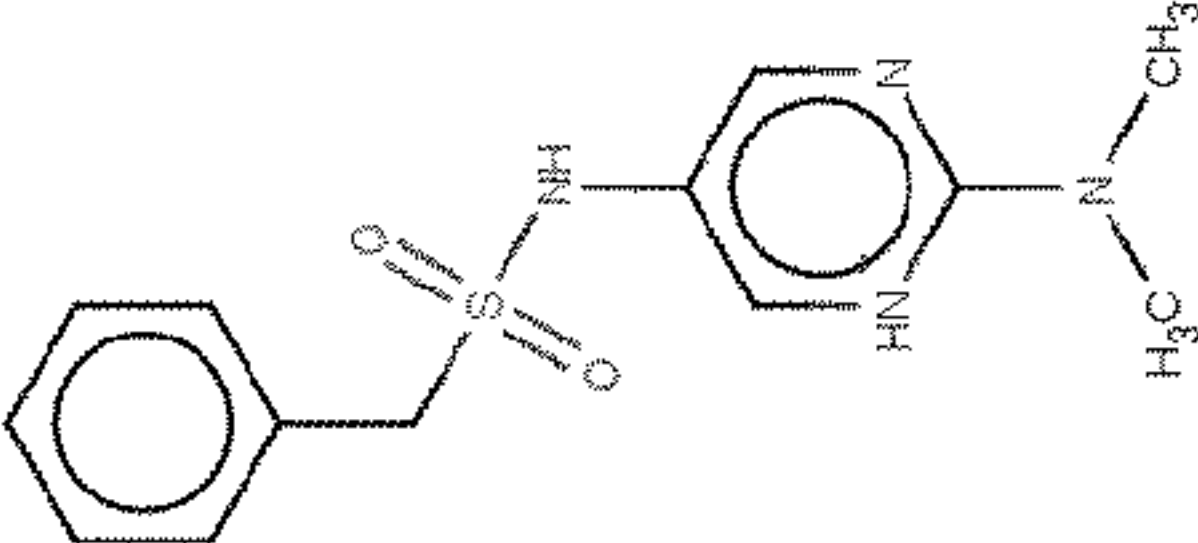
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 247		F5871-2878		F5871-2878_1 (1 of 2)
Compound 248		F5871-2883		F5871-2883_1 (1 of 2)
Compound 249		F5871-2885		F5871-2885_1_1 (1 of 2)

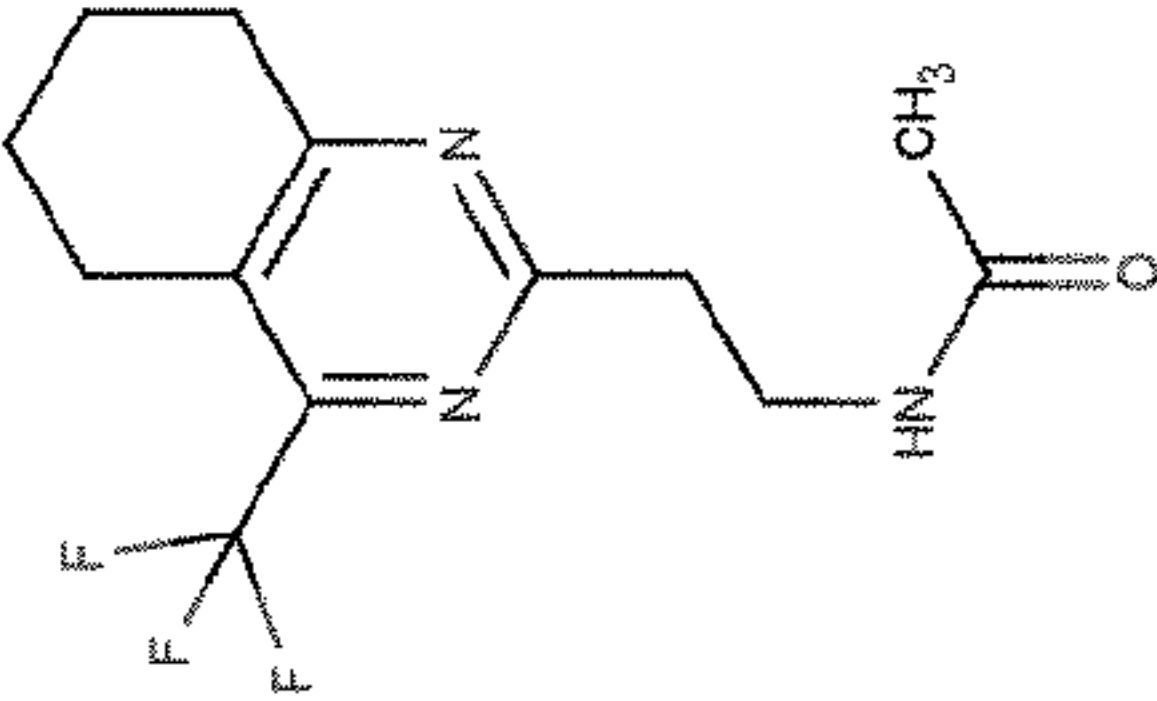
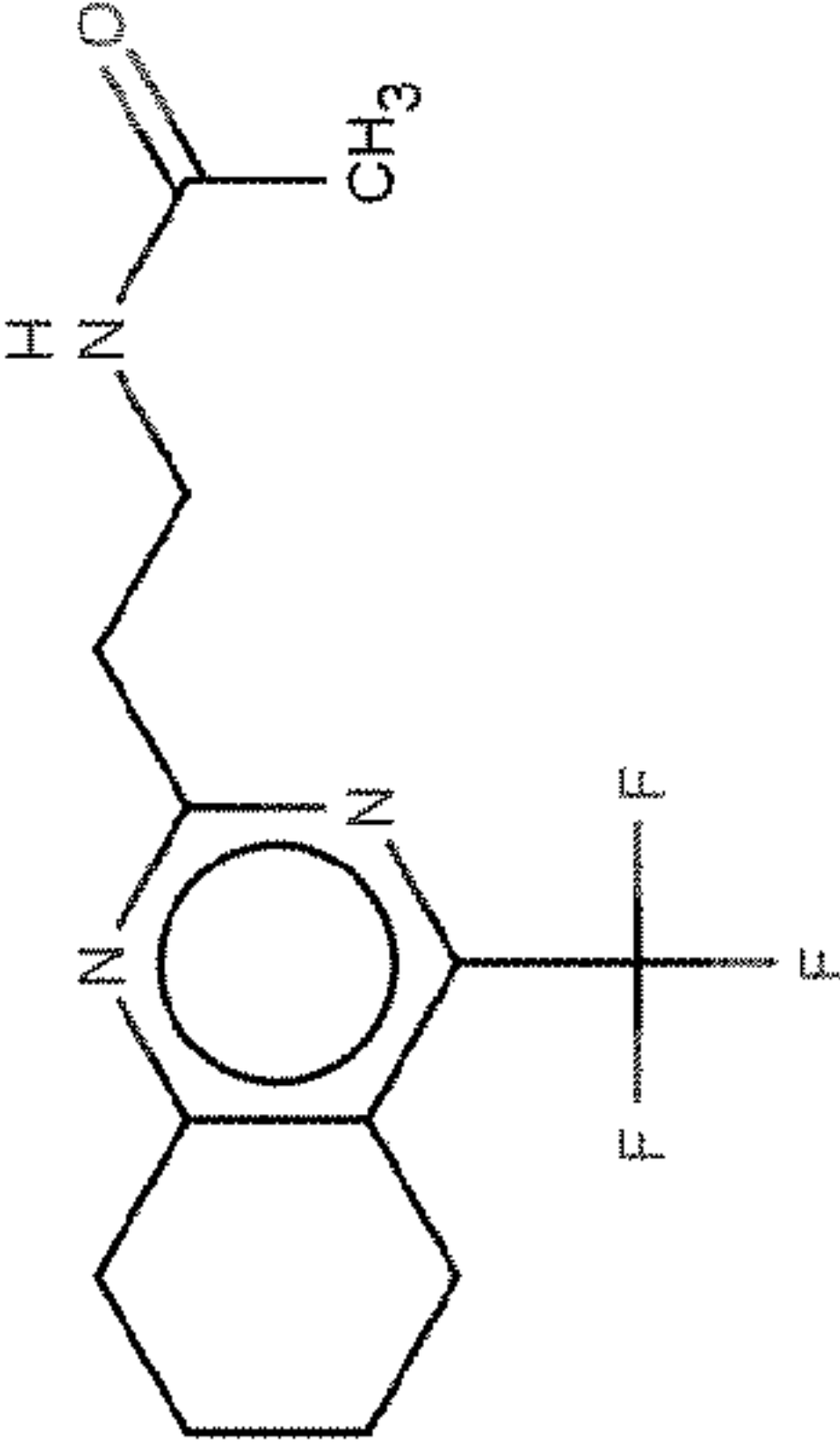
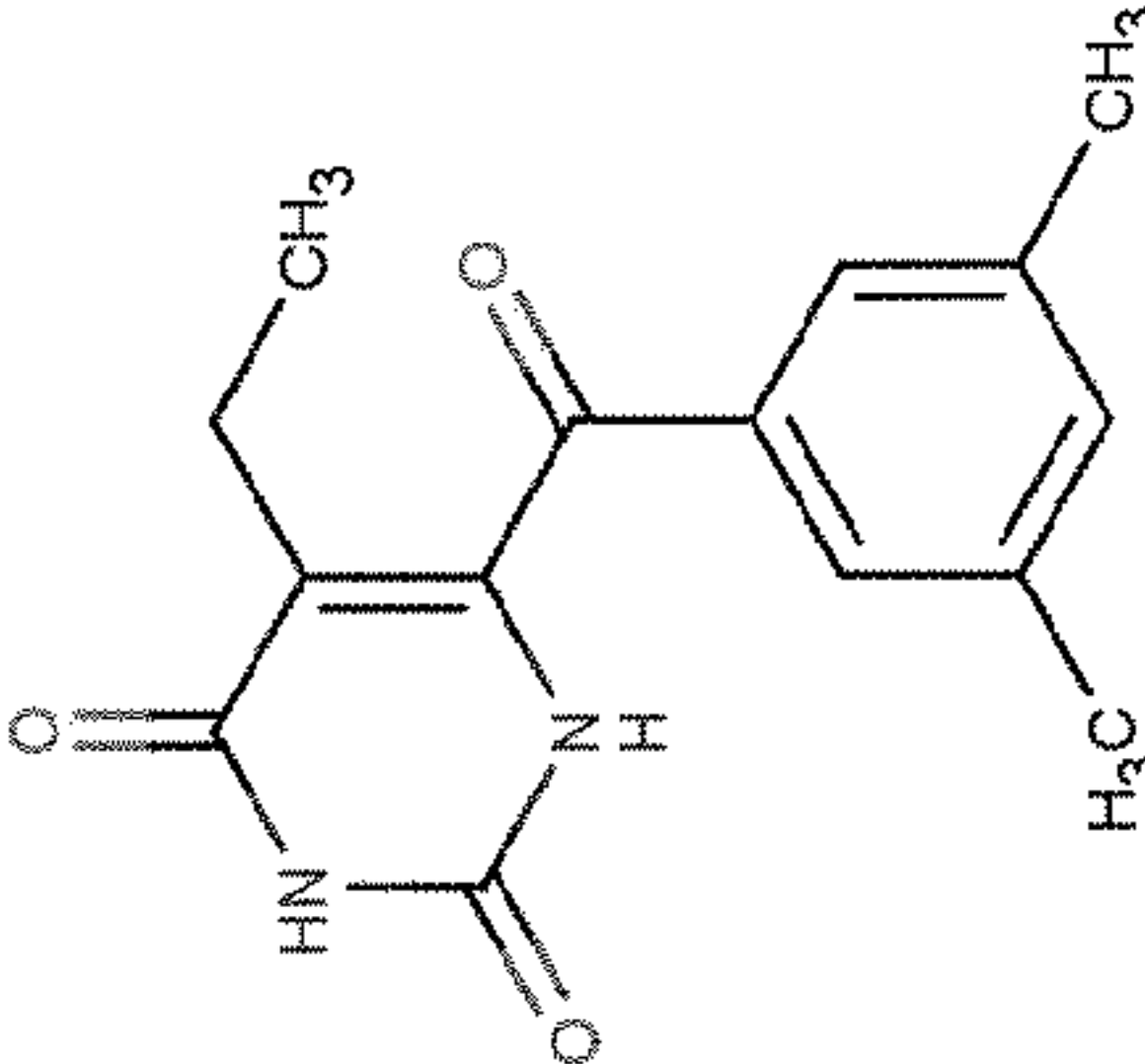
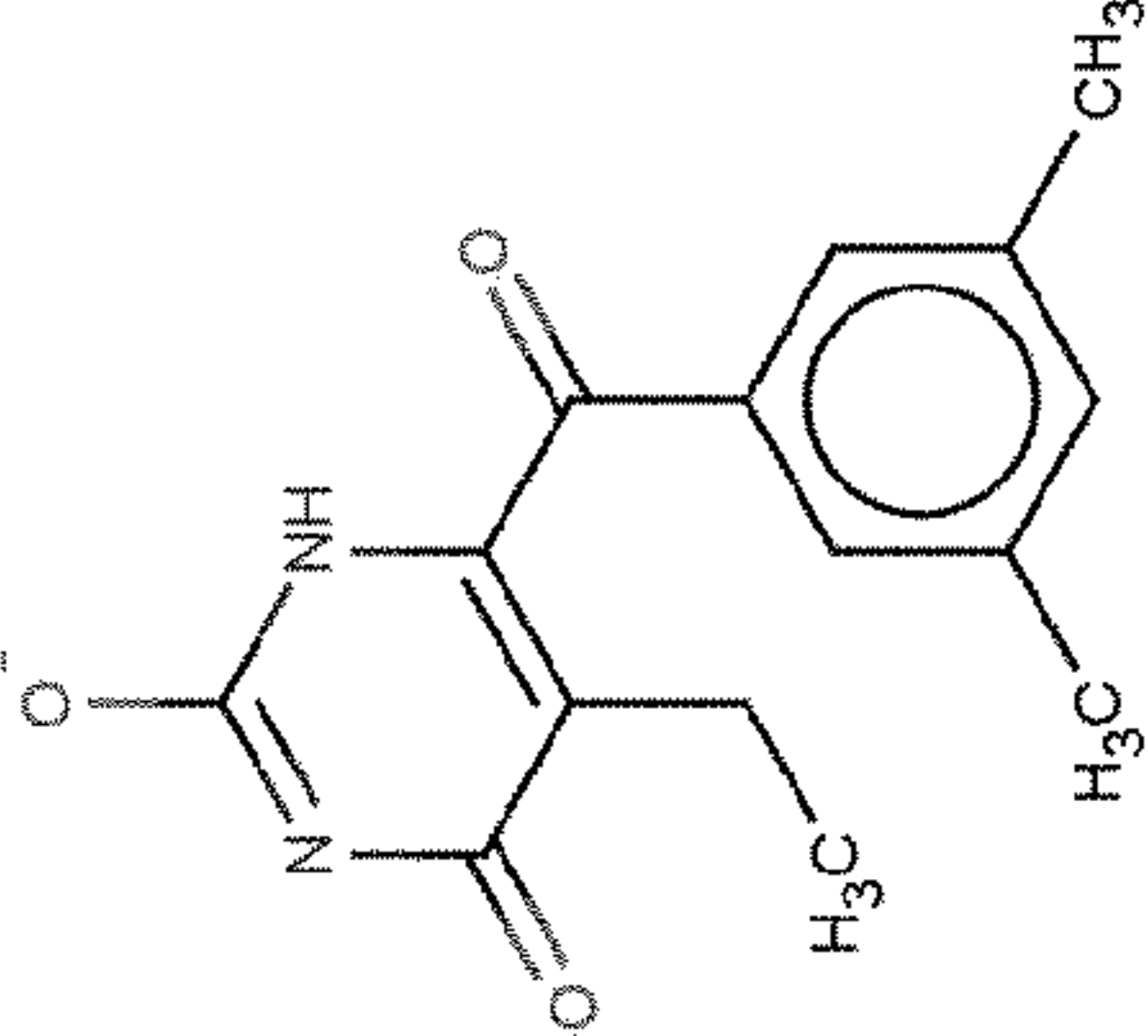
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 250		F5871-2898		F5871-2898_1 (1 of 2)
Compound 251		F5871-3594		F5871-3594_1_1 (1 of 2)
Compound 252		F5871-3603		F5871-3603_2_1 (1 of 3)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 253		F5871-3626		F5871-3626_1_1 (1 of 4)
Compound 254		F5871-3665		F5871-3665_3_1 (1 of 4)
Compound 255		F5871-4854		F5871-4854_1 (1 of 2)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 256		F5882-4189		F5882-4189_1_1 (1 of 8)
Compound 257		F5950-0041		F5950-0041_3_1 (1 of 4)
Compound 258		F5959-0024		F5959-0024_1_1 (1 of 7)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 259		F6037-0307		F6037-0307_3_1 (1 of 4)
Compound 260		F6064-0318		F6064-0318_3_1 (1 of 4)
Compound 261		F6064-0331		F6064-0331_1_1 (1 of 4)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 262		F6089-7863		F6089-7863_1_1 (1 of 2)
Compound 263		F6210-1183		F6210-1183_2 (1 of 2)
Compound 264		F6223-0039		F6223-0039_1_1 (1 of 2)

Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 265		F6286-0747		F6286-0747_1 (1 of 3)
Compound 266		F9995-0220		F9995-0220_22_1 (1 of 10)
Compound 267	$\text{CH}_3\text{OC}(\text{O})\text{CH}_2\text{C}(\text{O})\text{CH}_2\text{C}(\text{O})\text{CH}_3$	F9999-0001		
Compound 268	$\text{NH}_2\text{C}(\text{S})\text{NH}_2$	F9999-0002		

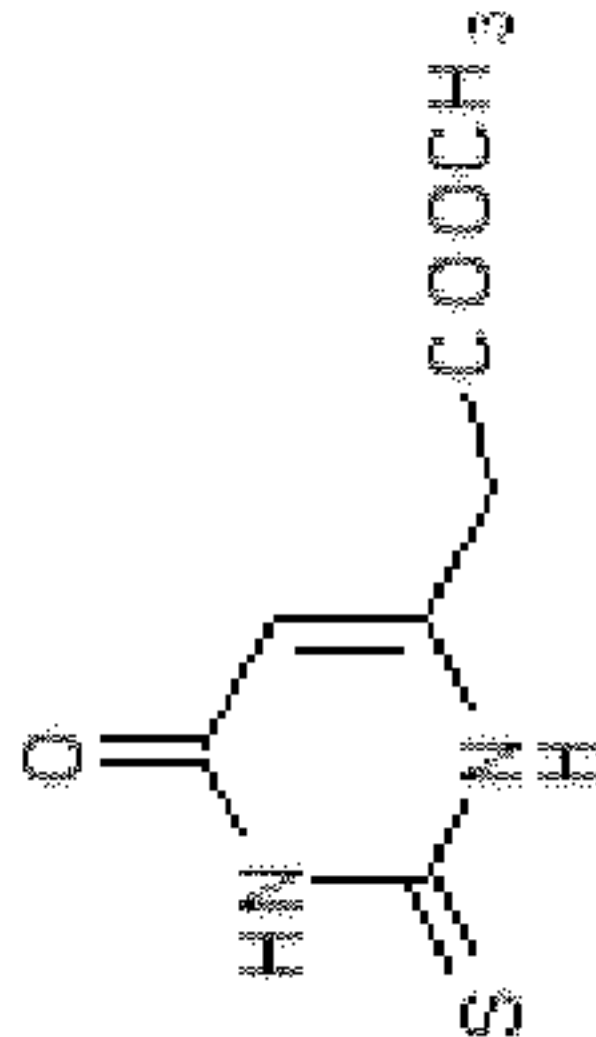
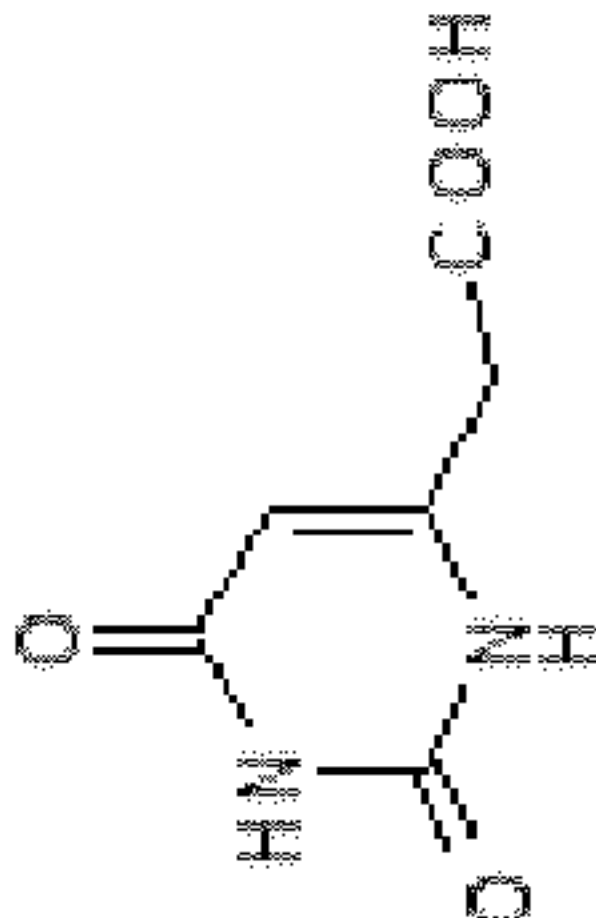
Compound x	Structure	No.	Tautomere/Stereoisomer	Reference of Variant
Compound 269		F9999-0003		
Compound 270		F9999-0004		

Table 1 presents compounds of the present invention as well as examples of tautomers and stereoisomers, respectively, the number of potential tautomer or stereoisomers is indicated in brackets in column 5, "reference of variant".

5

A chemical modification of compounds 1 to 270 is any modification that results in a compound characterized by a specific and effective interaction with or binding to the dimerization site of MIA. Chemical modifications are for example selected from the group consisting of H, alkyl, cycloalkyl, heterocycloalkyl, alkenyl, alkynyl, aryl, heteroaryl or alkyl, arylalkyl, cycloalkylalkyl, heterocycloalkylalkyl, heteroaryalkyl, and alkoxyalkyl, each of which is optionally substituted one, two or three times with independently selected non-polar or polar groups.

"Alkyl" as used herein alone or as part of another group, refers to a straight or branched chain hydrocarbon containing from 1 to 10 carbon atoms. Representative examples of alkyl include, but are not limited to, methyl, ethyl, n-propyl, iso-propyl, n-butyl, sec-butyl, iso-butyl, tert-butyl, n-pentyl, isopentyl, neopentyl, n-hexyl, 3-methylhexyl, 2,2-dimethylpentyl, 2,3-dimethylpentyl, n-heptyl, n-octyl, n-nonyl, n-decyl, and the like. "Lower alkyl" as used herein, is a subset of alkyl, in some embodiments preferred, and refers to a straight or branched chain hydrocarbon group containing from 1 to 4 carbon atoms. Representative examples of lower alkyl include, but are not limited to, methyl, ethyl, n-propyl, iso-propyl, n-butyl, iso-butyl, tert-butyl, and the like. The term "alkyl" or "loweralkyl" is intended to include both substituted and unsubstituted alkyl or loweralkyl unless otherwise indicated and these groups may be substituted with groups selected from halo (e.g., haloalkyl), alkyl, haloalkyl, alkenyl, alkynyl, cycloalkyl, cycloalkylalkyl, aryl, arylalkyl, heterocyclo, heterocycloalkyl, hydroxyl, alkoxy (thereby creating a polyalkoxy such as polyethylene glycol), alkenyloxy, alkynyloxy, haloalkoxy, cycloalkoxy, cycloalkylalkyloxy, aryloxy, arylalkyloxy, heterocycloxy, heterocyclolalkyloxy, mercapto, alkyl-S(0)m, haloalkyl-S(0)m, alkenyl-S(0)m, alkynyl-S(0)m, cycloalkyl-S(0)m, cycloalkylalkyl-S(0)m, aryl-S(0)m, arylalkyl-S(0)m, heterocyclo-S(0)m, heterocycloalkyl-S(0)m, amino, carboxy, alkylamino, alkenylamino, alkynylamino, haloalkylamino, cycloalkylamino, cycloalkylalkylamino, arylamino, arylalkylamino, heterocycloamino, heterocycloalkylamino, disubstituted-amino, acylamino, acyloxy, ester, amide, sulfonamide, urea, alkoxy acylamino, aminoacyloxy, nitro or cyano where m= 0, 1, 2 or 3.

"Alkenyl" as used herein alone or as part of another group, refers to a straight or branched chain hydrocarbon containing from 1 to 10 carbon atoms (or in lower alkenyl 1 to 4 carbon atoms) which include 1 to 4 double bonds in the normal chain.

5 Representative examples of alkenyl include, but are not limited to, vinyl, 2-propenyl, 3-butenyl, 2-butenyl, 4-pentenyl, 3-pentenyl, 2-hexenyl, 3-hexenyl, 2,4-heptadiene, and the like. The term "alkenyl" or "lower alkenyl" is intended to include both substituted and unsubstituted alkenyl or loweralkenyl unless otherwise indicated and these groups may be substituted with groups as described in connection with alkyl and loweralkyl
10 above.

"Alkynyl" as used herein alone or as part of another group, refers to a straight or branched chain hydrocarbon containing from 1 to 10 carbon atoms (or in loweralkynyl 1 to 4 carbon atoms) which include 1 triple bond in the normal chain. Representative
15 examples of alkynyl include, but are not limited to, 2-propynyl, 3-butylnyl, 2-butylnyl, 4-pentylnyl, 3-pentylnyl, and the like. The term "alkynyl" or "loweralkynyl" is intended to include both substituted and unsubstituted alkynyl or loweralkynyl unless otherwise indicated and these groups may be substituted with the same groups as set forth in connection with alkyl and loweralkyl above.

20 "Cycloalkyl" as used herein alone or as part of another group, refers to a saturated or partially unsaturated cyclic hydrocarbon group containing from 3, 4 or 5 to 6, 7 or 8 carbons (which carbons may be replaced in a heterocyclic group as discussed below). Representative examples of cycloalkyl include, cyclopropyl, cyclobutyl, cyclopentyl,
25 cyclohexyl, cycloheptyl, and cyclooctyl. These rings may be optionally substituted with additional substituents as described herein such as halo or lower alkyl. The term "cycloalkyl" is generic and intended to include heterocyclic groups as discussed below unless specified otherwise.

30 "Heterocyclic group" or "heterocyclo" as used herein alone or as part of another group, refers to an aliphatic (e.g., fully or partially saturated heterocyclo) or aromatic (e.g., heteroaryl) monocyclic- or a bicyclic-ring system. Monocyclic ring systems are exemplified by any 5 or 6 membered ring containing 1, 2, 3, or 4 heteroatoms independently selected from oxygen, nitrogen and sulfur. The 5 membered ring has from
35 0-2 double bonds and the 6 membered ring has from 0-3 double bonds. Representative

examples of monocyclic ring systems include, but are not limited to, azetidine, azepine, aziridine, diazepine, 1,3-dioxolane, dioxane, dithiane, furan, imidazole, imidazoline, imidazolidine, isothiazole, isothiazoline, isothiazolidine, isoxazole, isoxazoline, isoxazolidine, morpholine, oxadiazole, oxadiazoline, oxadiazolidine, oxazole, oxazoline, oxazolidine, piperazine, piperidine, pyran, pyrazine, pyrazole, pyrazoline, pyrazolidine, pyridine, pyrimidine, pyridazine, pyrrole, pyrroline, pyrrolidine, tetrahydrofuran, tetrahydrothiophene, tetrazine, tetrazole, thiadiazole, thiadiazoline, thiadiazolidine, thiazole, thiazoline, thiazolidine, thiophene, thiomorpholine, thiomorpholine sulfone, thiopyran, triazine, triazole, trithiane, and the like. Bicyclic ring systems are exemplified by any of the above monocyclic ring systems fused to an aryl group as defined herein, a cycloalkyl group as defined herein, or another monocyclic ring system as defined herein. Representative examples of bicyclic ring systems include but are not limited to, for example, benzimidazole, benzothiazole, benzothiadiazole, benzothiophene, benzoxadiazole, benzoxazole, benzofuran, benzopyran, benzothiopyran, benzodioxine, 1,3- benzodioxole, cinnoline, indazole, indole, indoline, indolizine, naphthyridine, isobenzofuran, isobenzothiophene, isoindole, isoindoline, isoquinoline, phthalazine, purine, pyranopyridine, quinoline, quinolizine, quinoxaline, quinazoline, tetrahydroisoquinoline, tetrahydroquinoline, thiopyranopyridine, and the like. These rings include quaternized derivatives thereof and may be optionally substituted with groups selected from halo, alkyl, haloalkyl, alkenyl, alkynyl, cycloalkyl, cycloalkylalkyl, aryl, arylalkyl, heterocyclo, heterocycloalkyl, hydroxyl, alkoxy, alkenyloxy, alkynyloxy, haloalkoxy, cycloalkoxy, cycloalkylalkyloxy, aryloxy, arylalkyloxy, heterocycloxy, heterocyclolalkyloxy, mercapto, alkyl-S(O)_m, haloalkyl-S(O)_m, alkenyl-S(O)_m, alkynyl-S(O)_m, cycloalkyl-S(O)_m, cycloalkylalkyl-S(O)_m, aryl-S(O)_m, arylalkyl-S(O)_m, heterocyclo-S(O)_m, heterocycloalkyl-S(O)_m, amino, alkylamino, alkenylamino, alkynylamino, haloalkylamino, cycloalkylamino, cycloalkylalkylamino, arylamino, arylalkylamino, heterocycloamino, heterocycloalkylamino, disubstituted-amino, acylamino, acyloxy, ester, amide, sulfonamide, urea, alkoxyacylamino, aminoacyloxy, nitro or cyano where m = 0, 1, 2 or 3.

"Aryl" as used herein alone or as part of another group, refers to a monocyclic carbocyclic ring system or a bicyclic carbocyclic fused ring system .having one or more aromatic rings. Representative examples of aryl include, azulenyl, indanyl, indenyl, naphthyl, phenyl, tetrahydronaphthyl, and the like. The term "aryl" is intended to include both substituted and unsubstituted aryl unless otherwise indicated and these groups may be

substituted with the same groups as set forth in connection with alkyl and loweralkyl above.

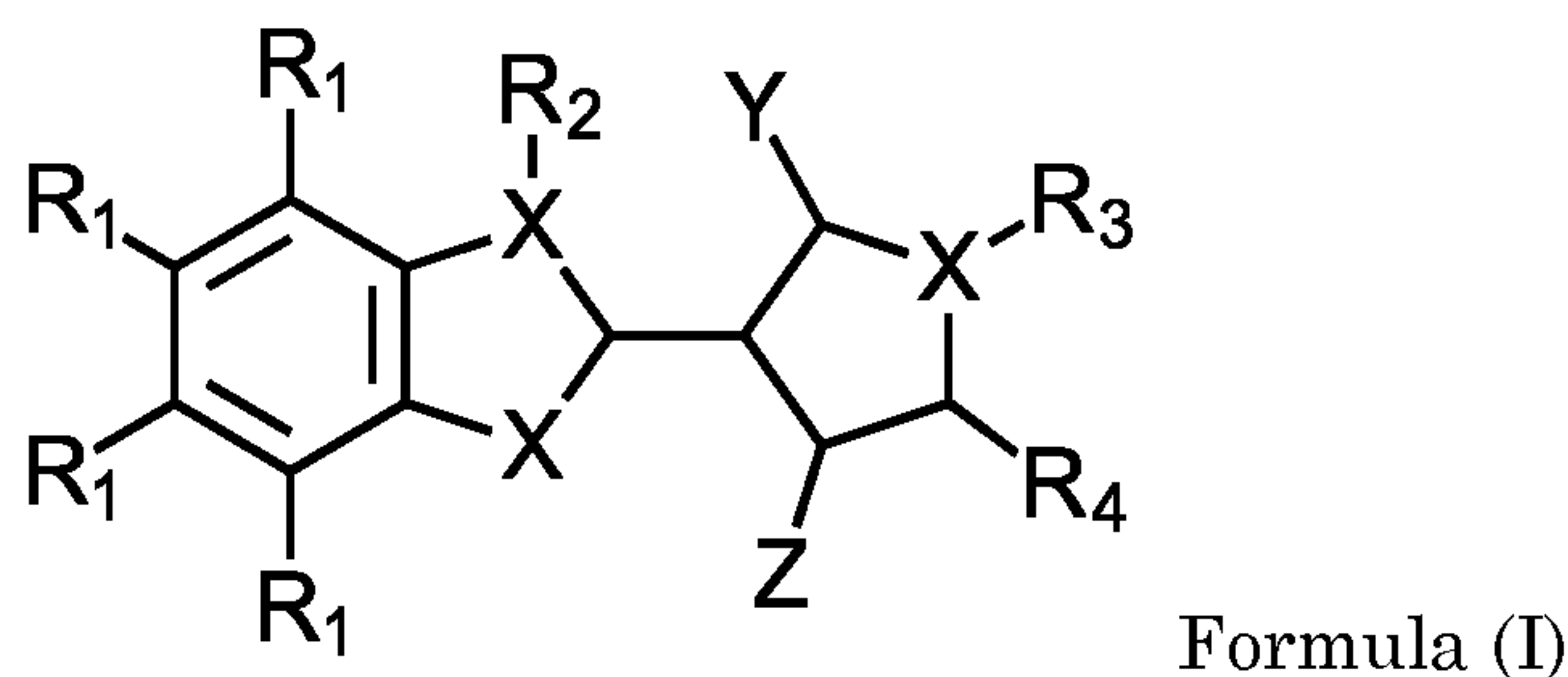
"Arylalkyl" as used herein alone or as part of another group, refers to an aryl group, as defined herein, appended to the parent molecular moiety through an alkyl group, as defined herein. Representative examples of arylalkyl include, but are not limited to, benzyl, 2-phenylethyl, 3-phenylpropyl, 2-naphth-2-ylethyl, and the like.

"Heteroaryl" as used herein is as described in connection with heterocyclo above.

"Alkoxy" as used herein alone or as part of another group, refers to an alkyl or loweralkyl group, as defined herein (and thus including substituted versions such as polyalkoxy), appended to the parent molecular moiety through an oxy group, -O-. Representative examples of alkoxy include, but are not limited to, methoxy, ethoxy, propoxy, 2-propoxy, butoxy, tert-butoxy, pentyloxy, hexyloxy and the like.

"Halo" as used herein refers to any suitable halogen, including -F, -Cl, -Br, and -I.

Compound 1 is represented by the following general Formula (I)



Formula (I)

wherein,

R₁, R₂ and R₄ represent, independent of each other, hydrogen, halogens, alkyls, alkenyls, alkynyls, cycloalkyls or aryls;

R₃ represents a linear n₁-7 alkyl, a branched n₃-7 alkyl, a linear n₁-7 alkenyl, a branched n₃-7 alkenyl, a linear n₁-7 alkynyl, a branched n₃-7 alkynyl, a n₅-7 cycloalkyl, a n₅-7 cycloalkenyl, a n₅-7 cycloalkynyl, a n₅-7 aryl;

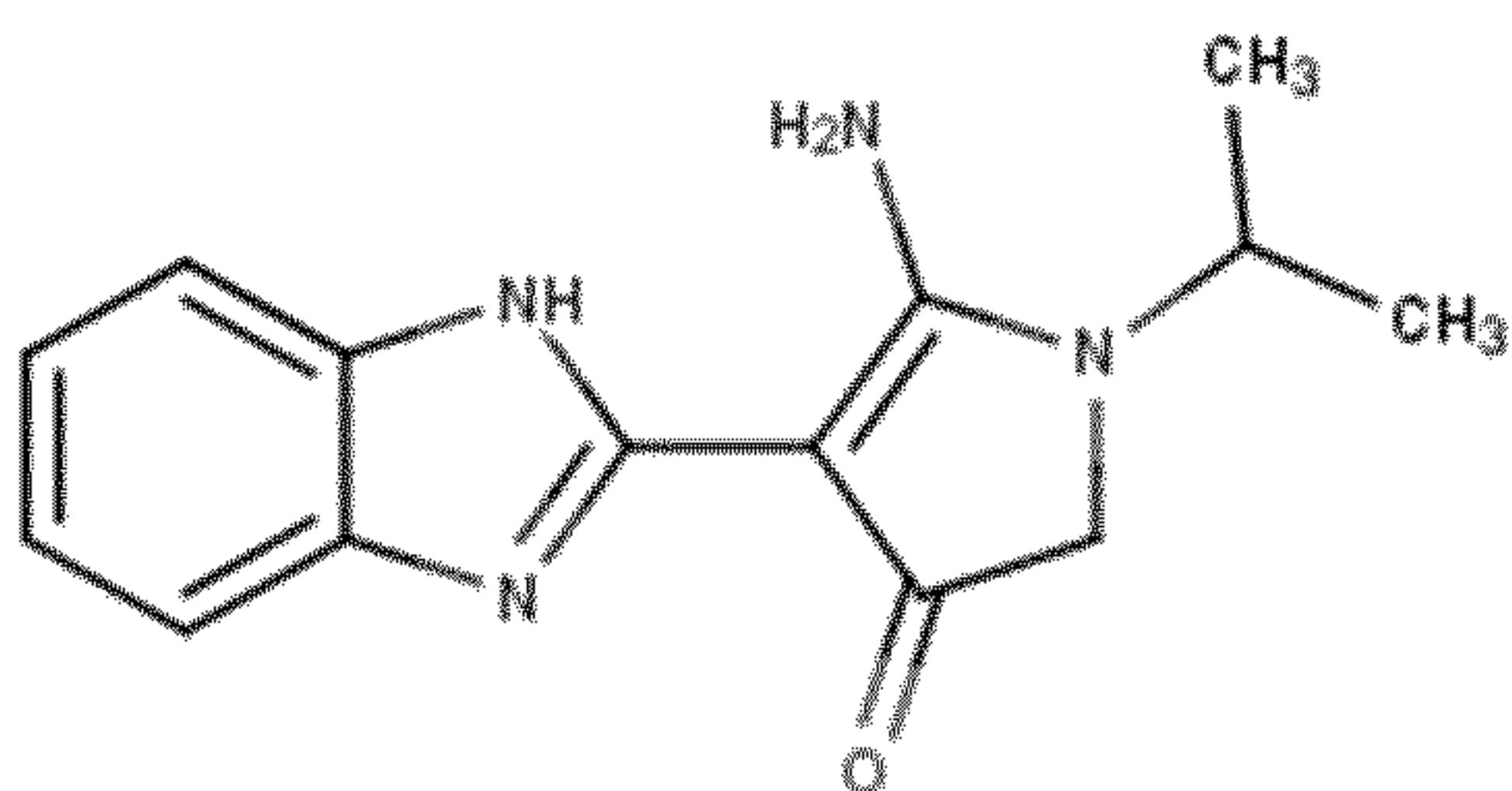
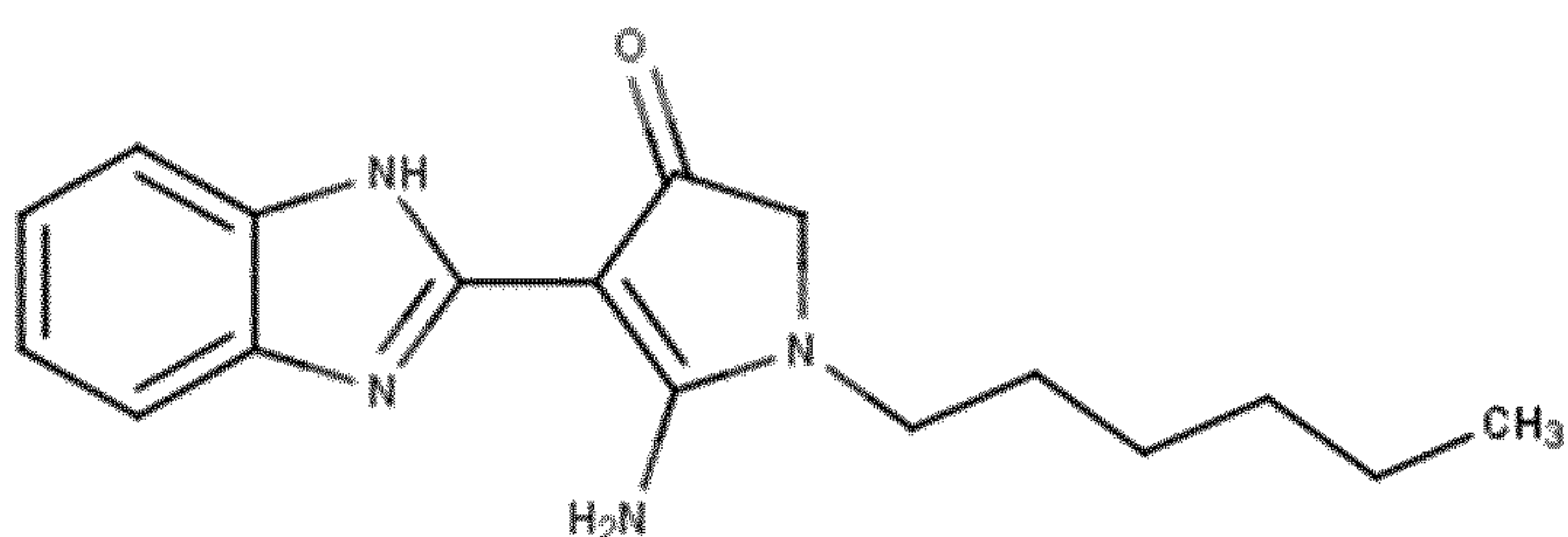
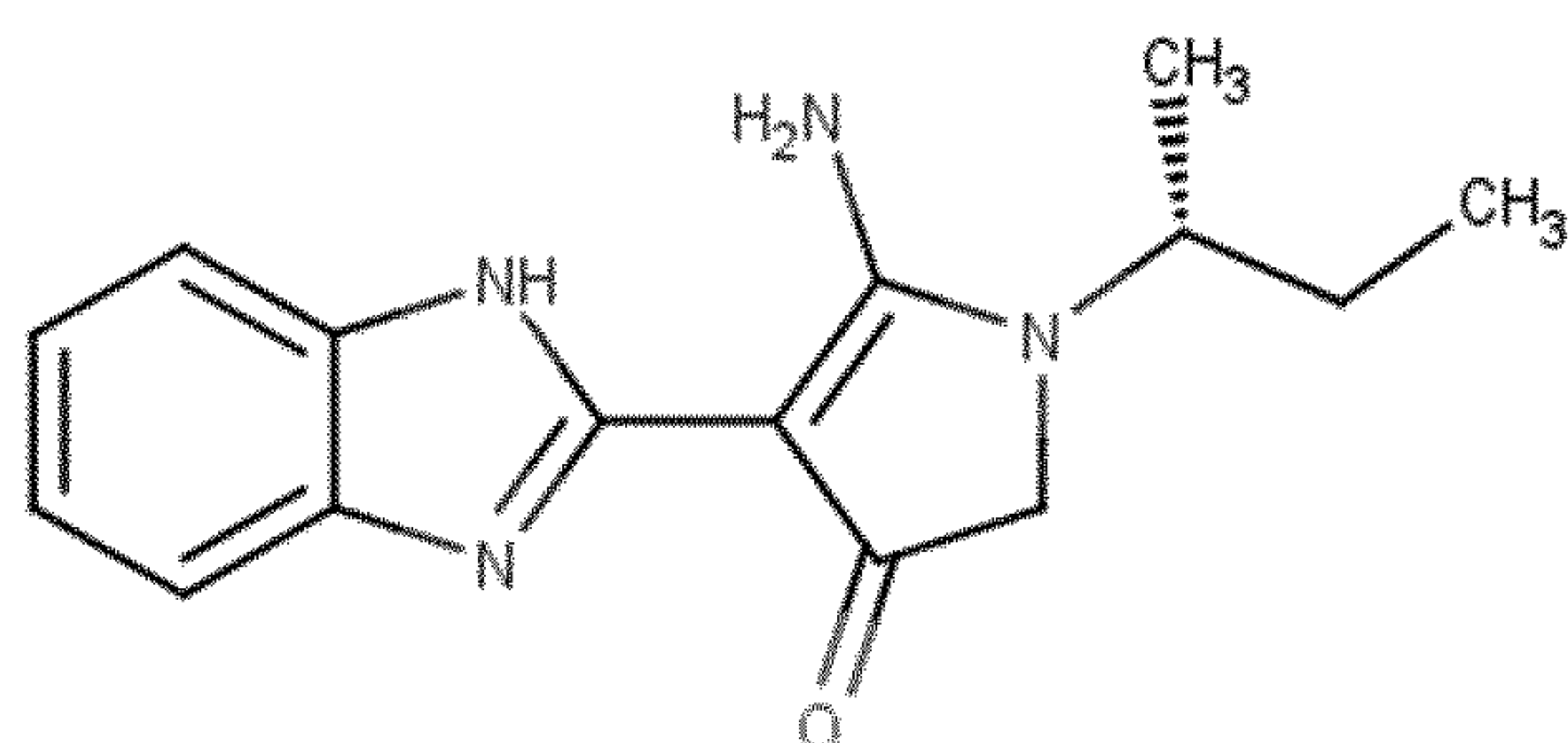
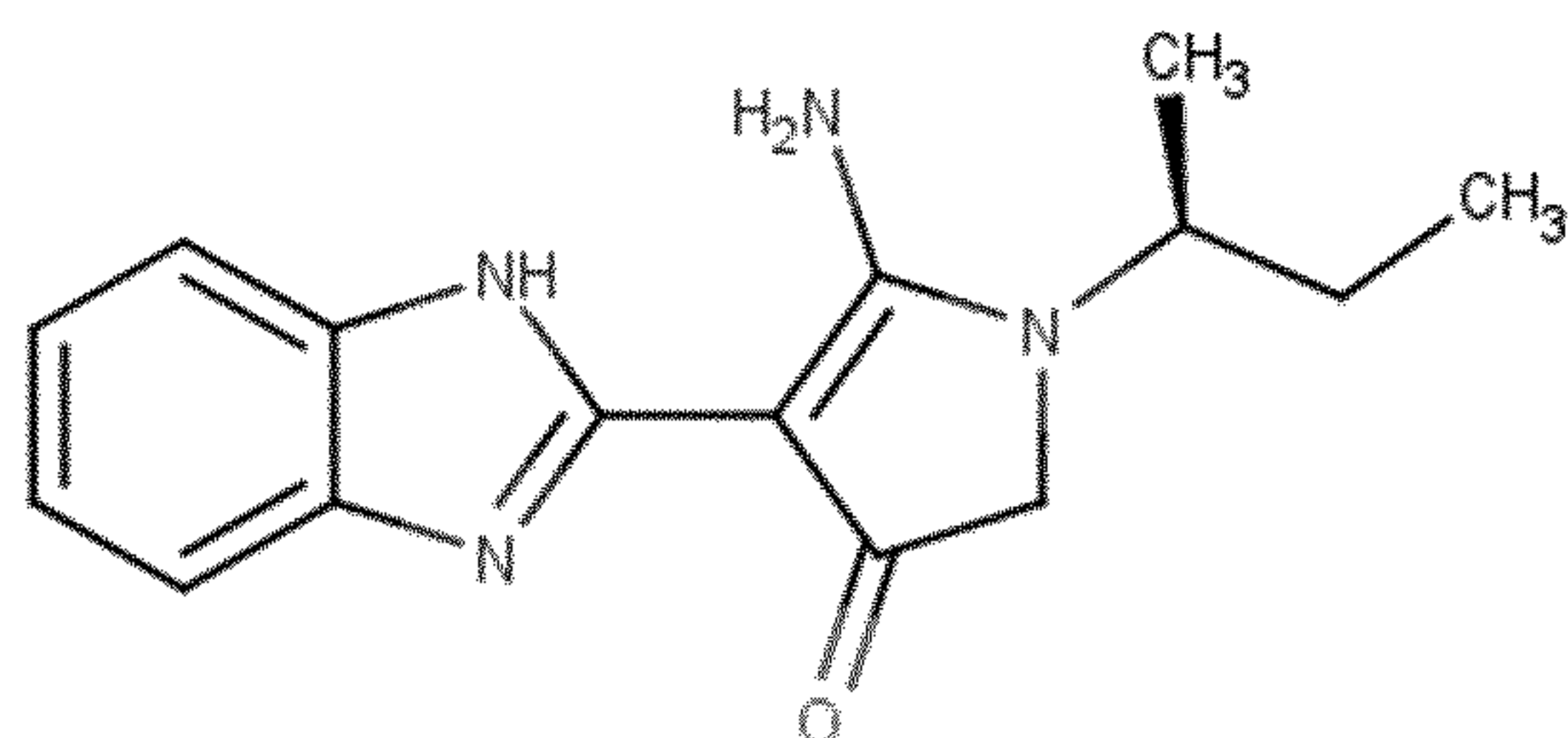
X represents, independent of each other, a substituted or unsubstituted heteroatom selected from N and S, wherein the heteroatom may form a double bond with a neighboring carbon atom;

Y represents a substituted or unsubstituted amine which may form a double bond with a neighboring carbon atom; and

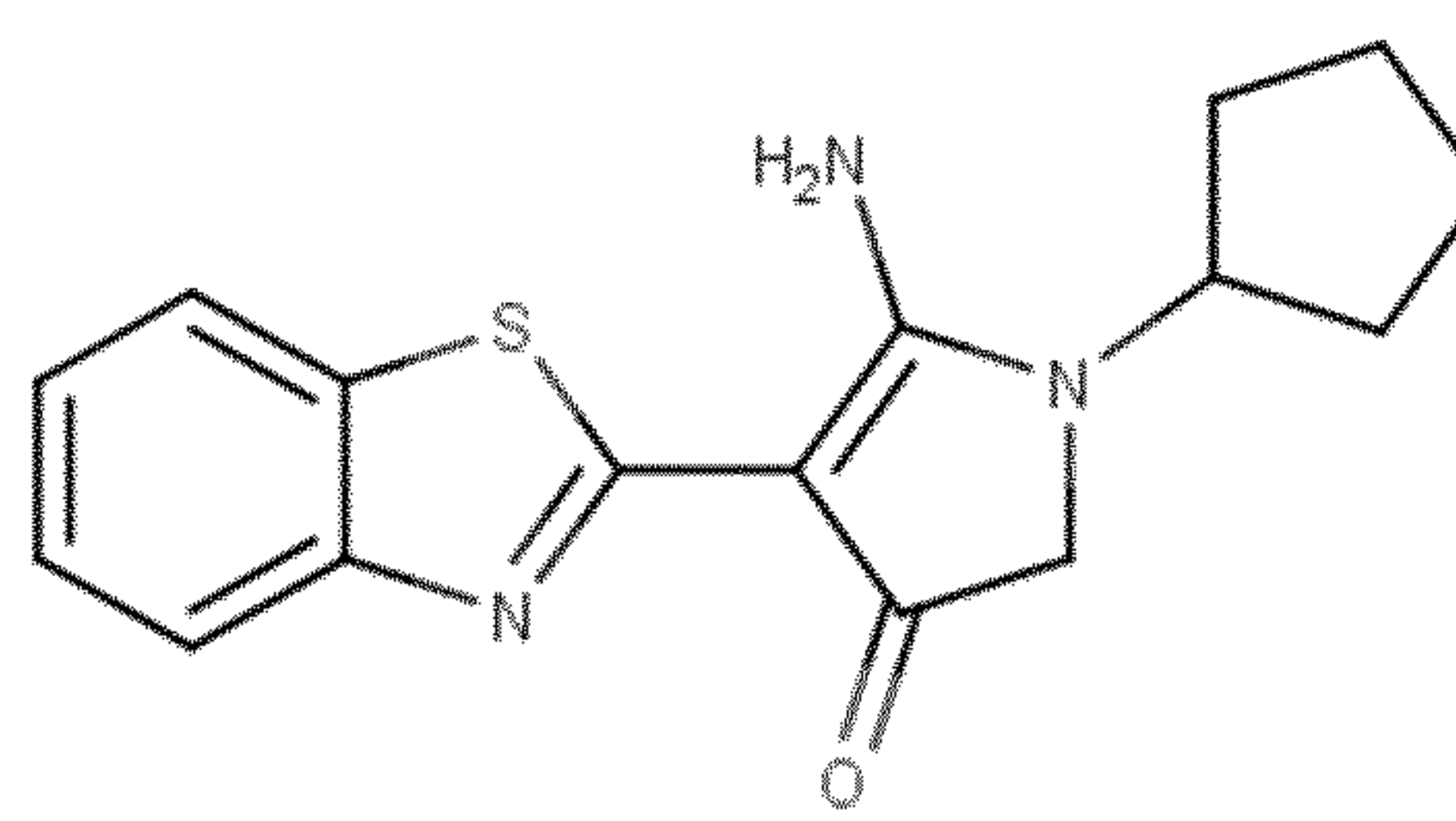
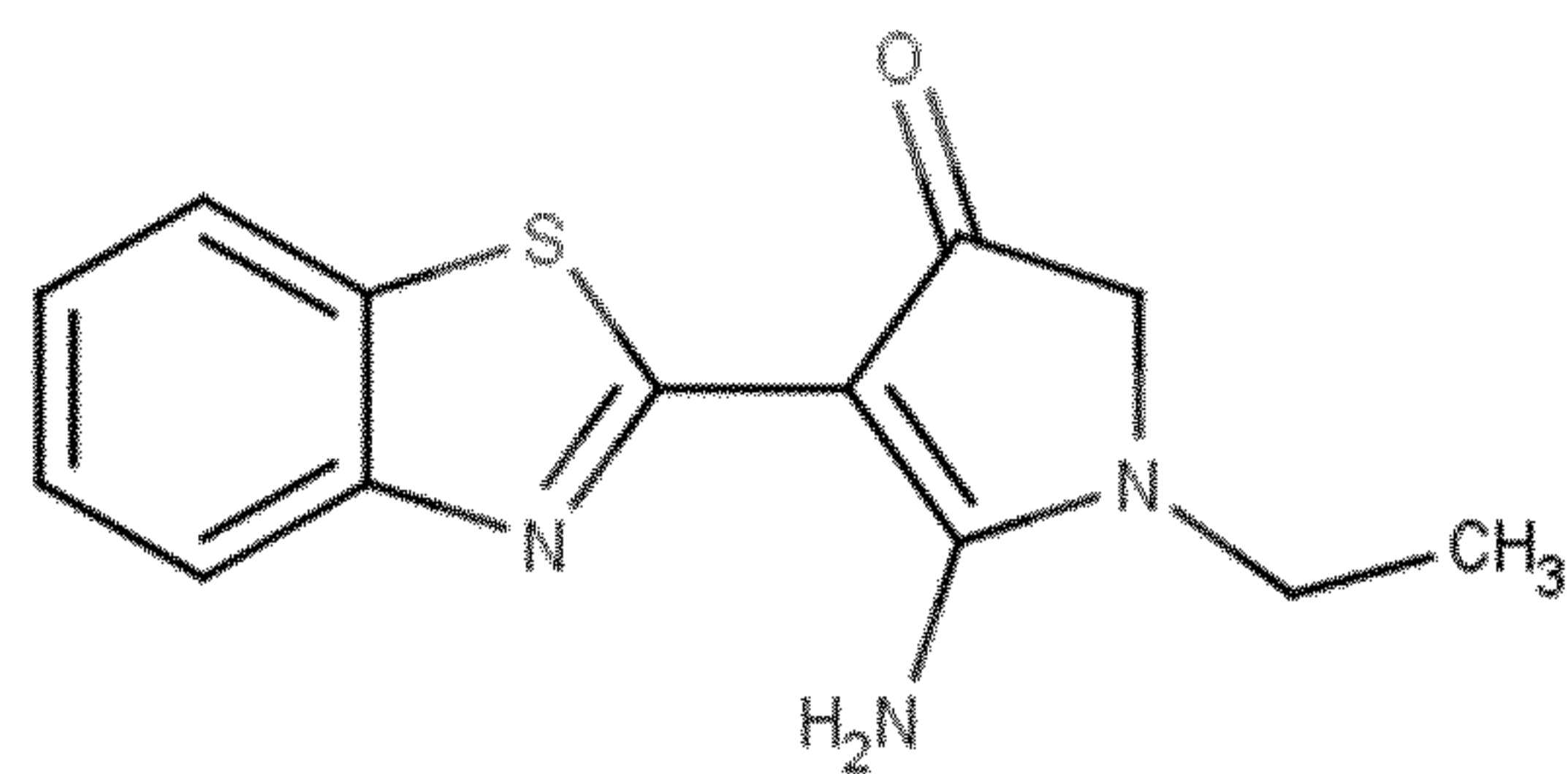
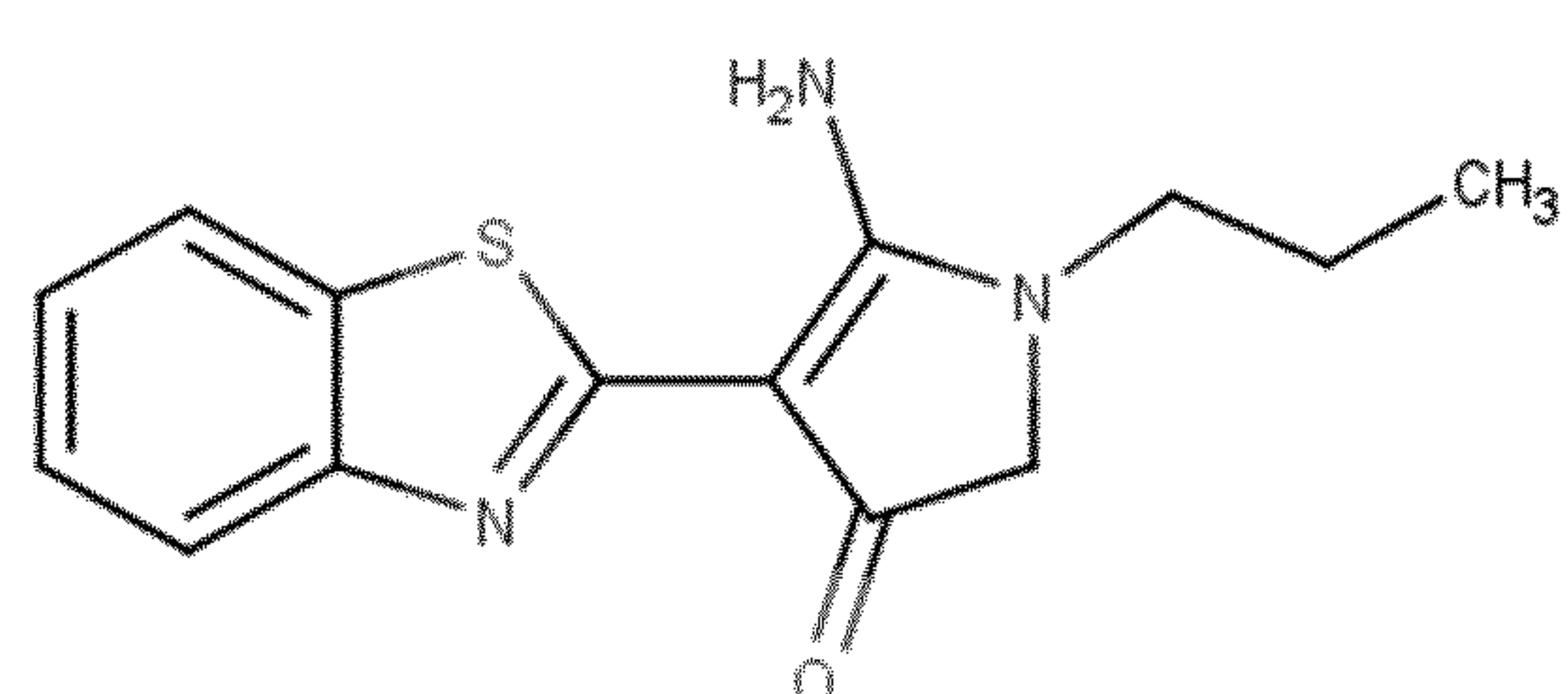
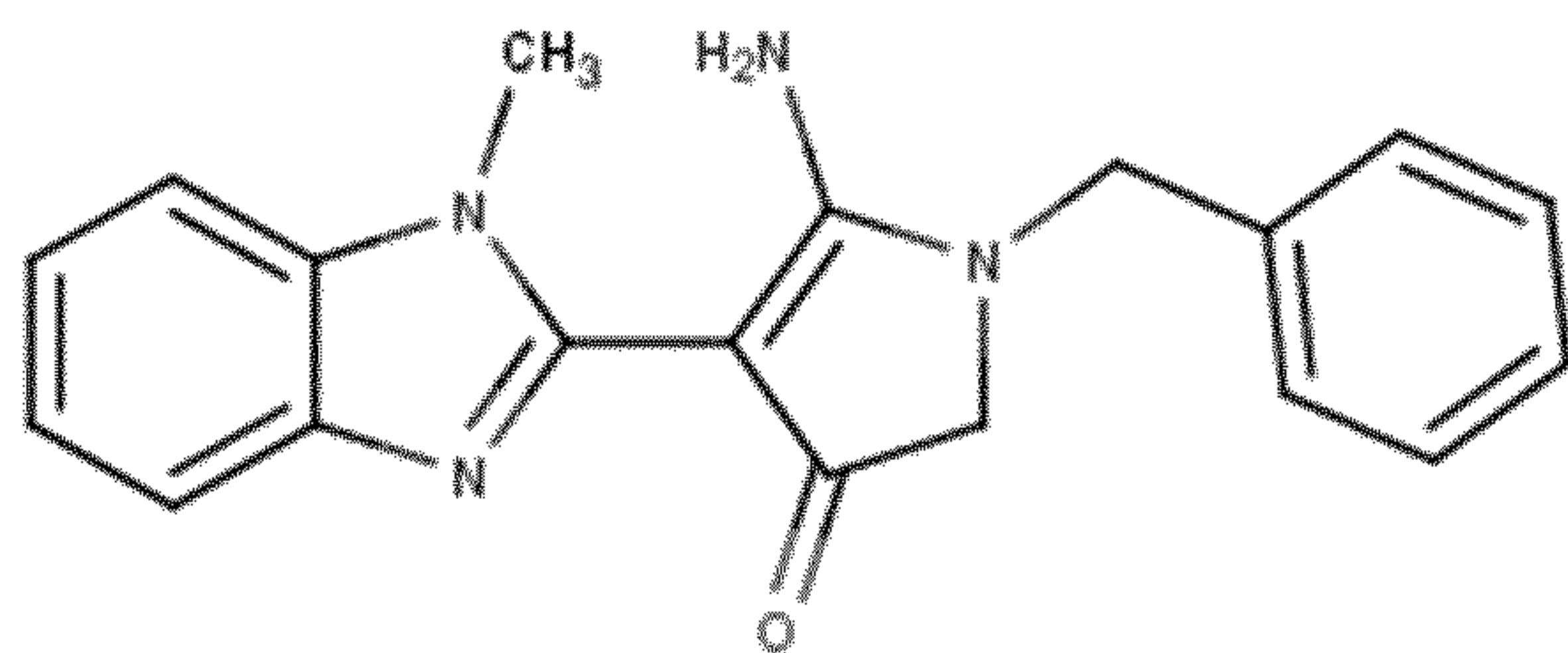
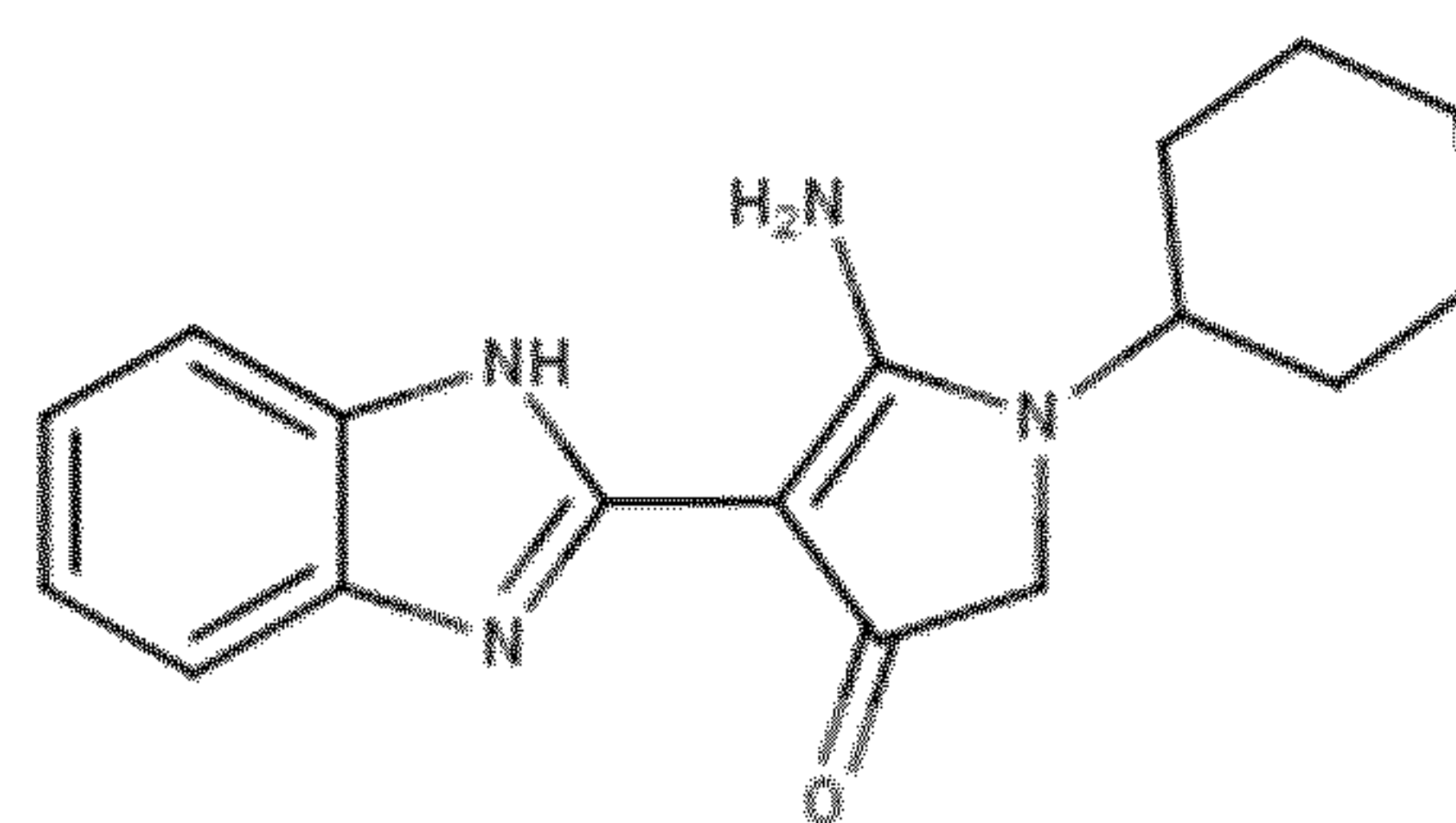
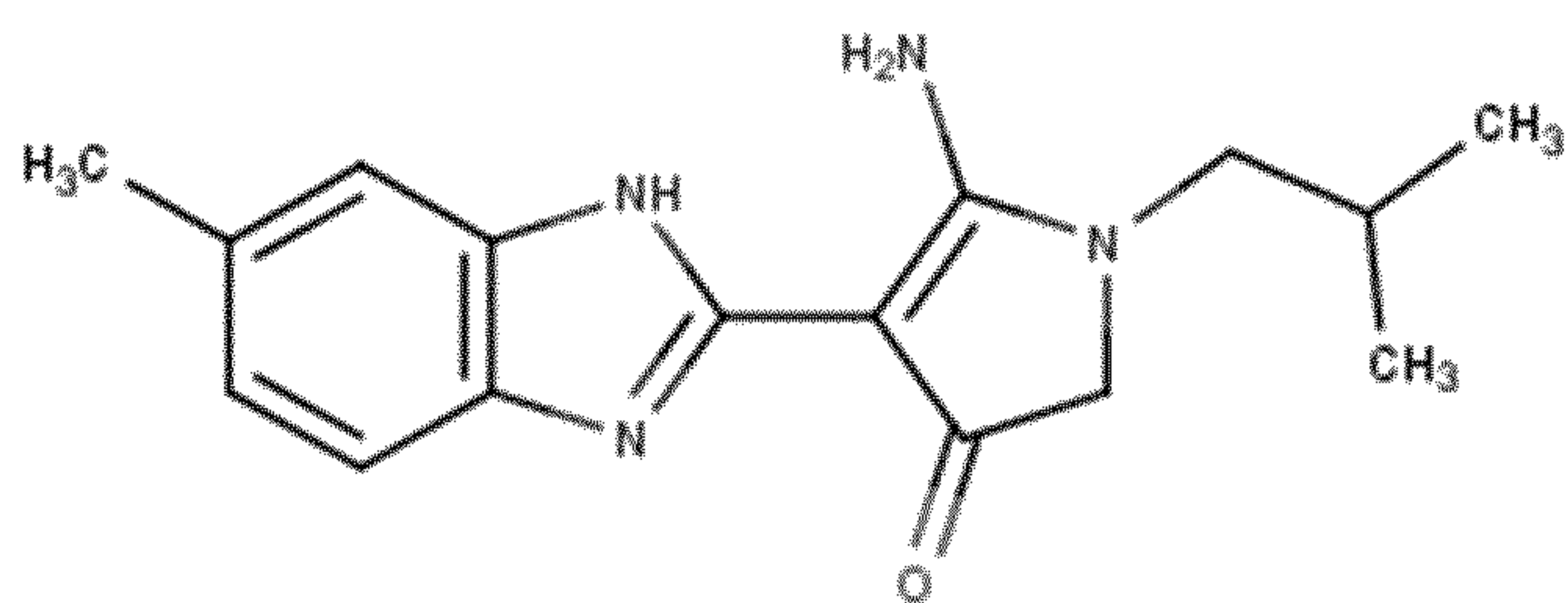
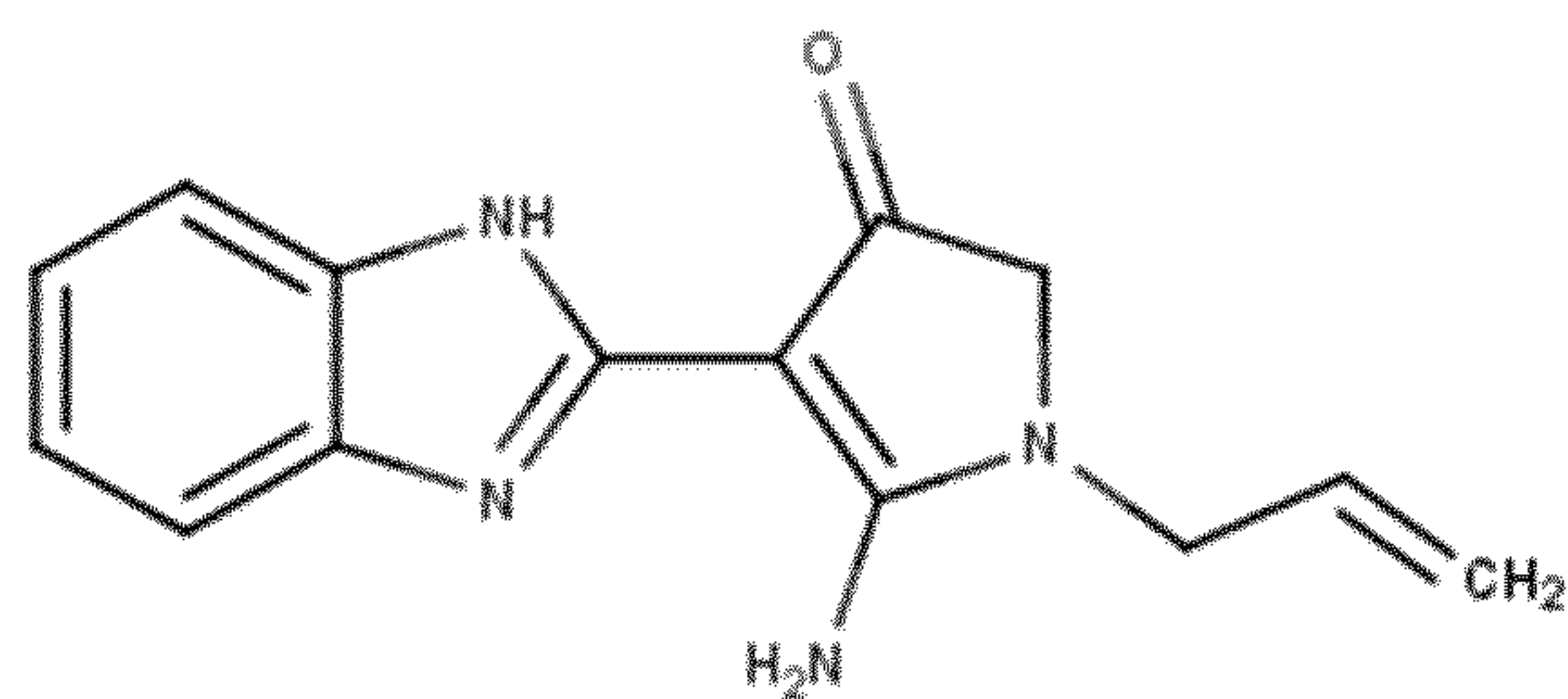
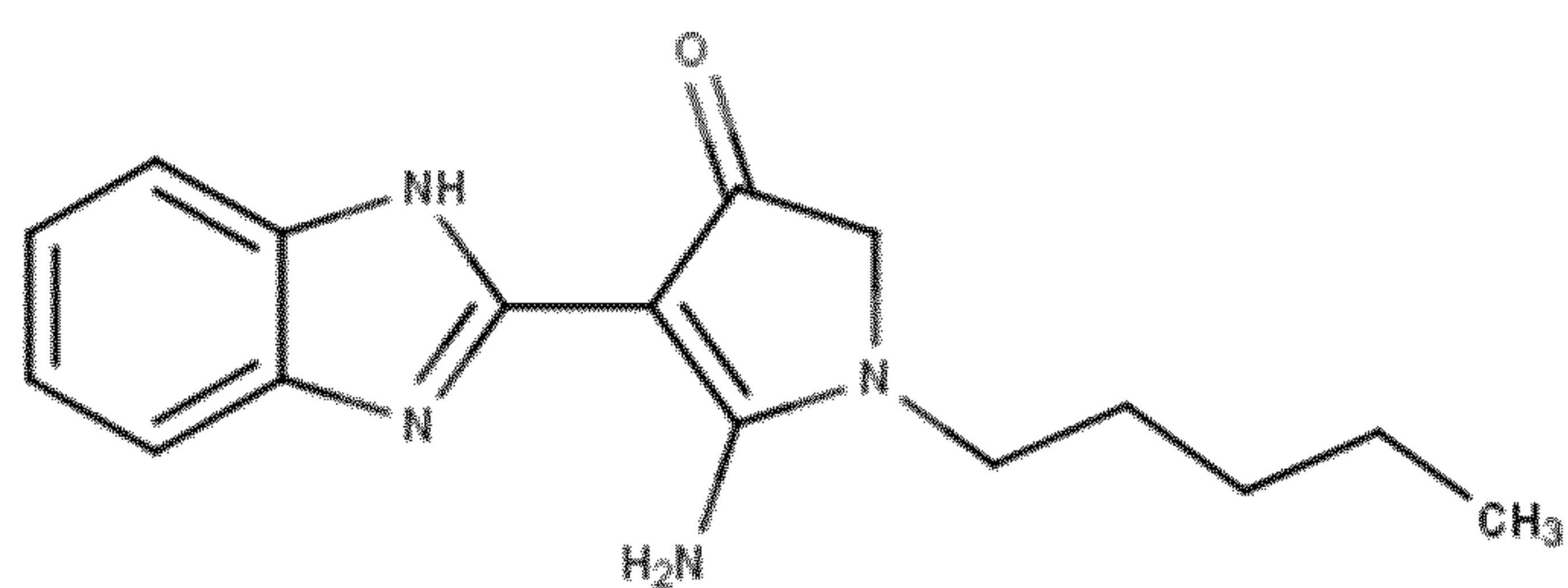
Z represents a halogen or a substituted or unsubstituted heteroatom selected from O, N and S, wherein the heteroatom may form a double bond with a neighboring carbon atom.

Compound 1 comprises for example the following modifications:

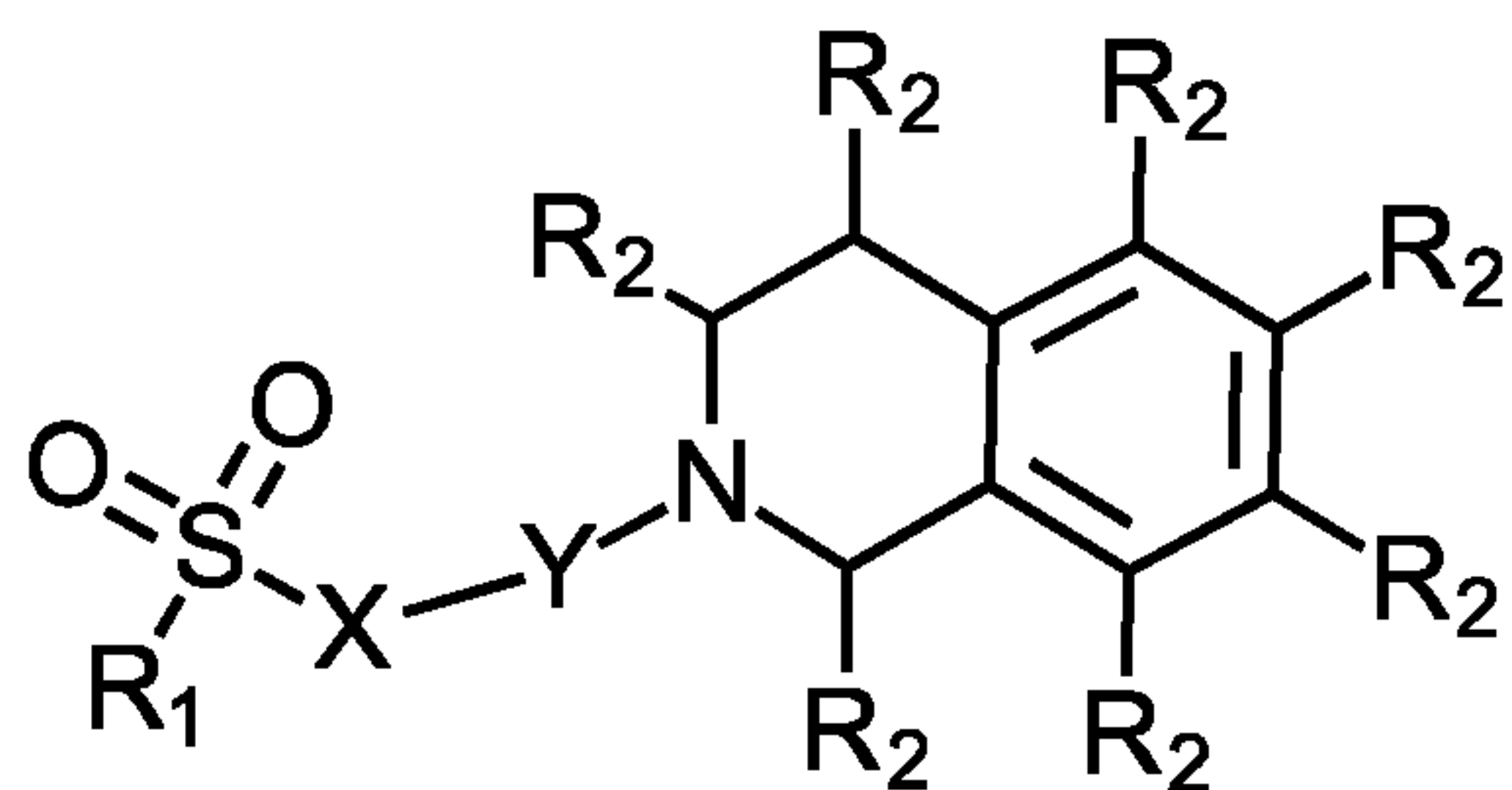
5



10



Compound 2 is represented by the following general Formula (II)



Formula (II)

wherein,

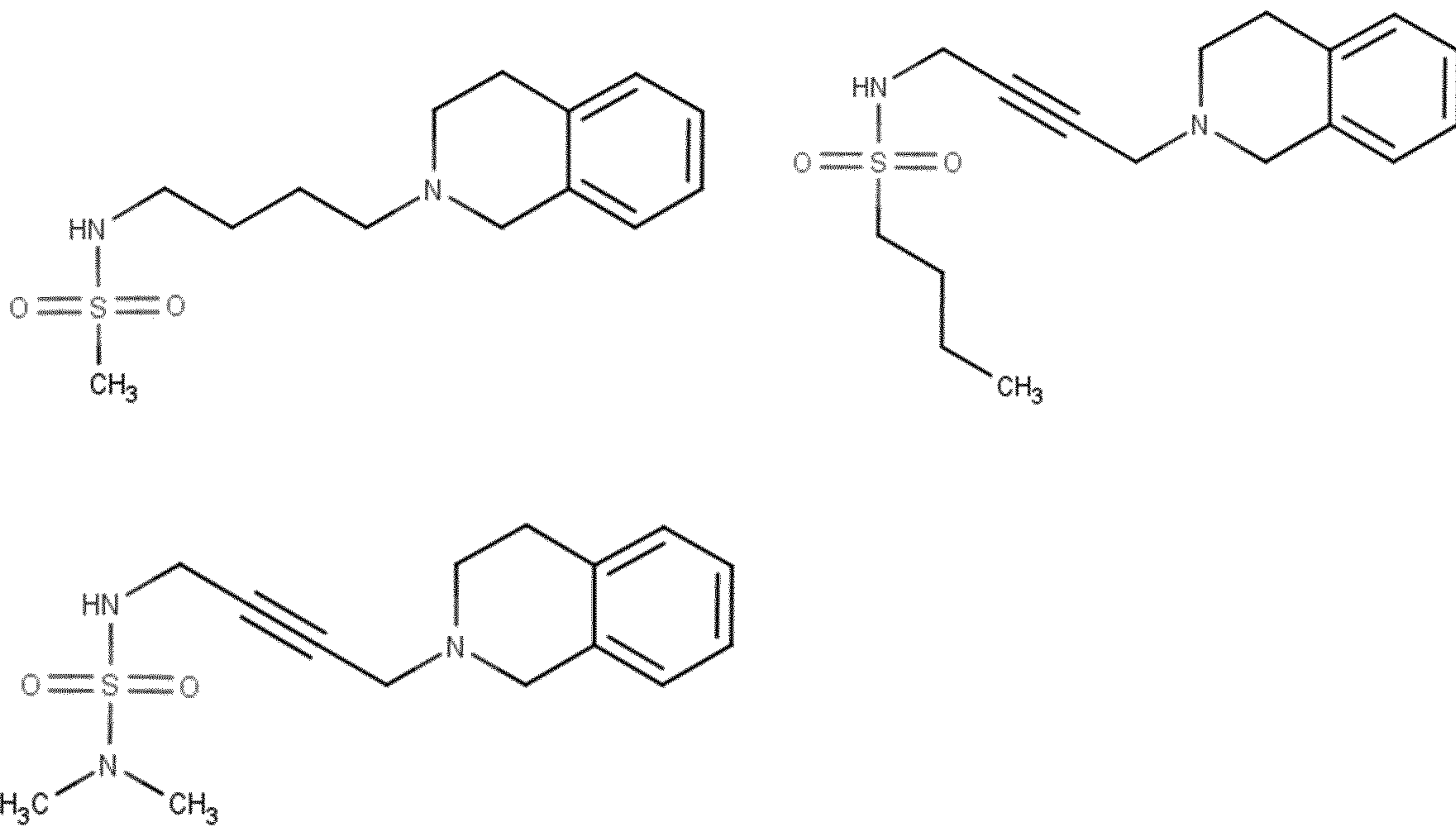
R₁ represents a hydrogen, alkyl, alkenyl, alkynyl, amin, aminoalkyl, aminoalkenyl, halogens, alkyls, alkenyls, alkynyls, cycloalkyls or aryl;

5 R₂ represents, independent from each other, a hydrogen, alkyl, alkenyl, or alkynyl;

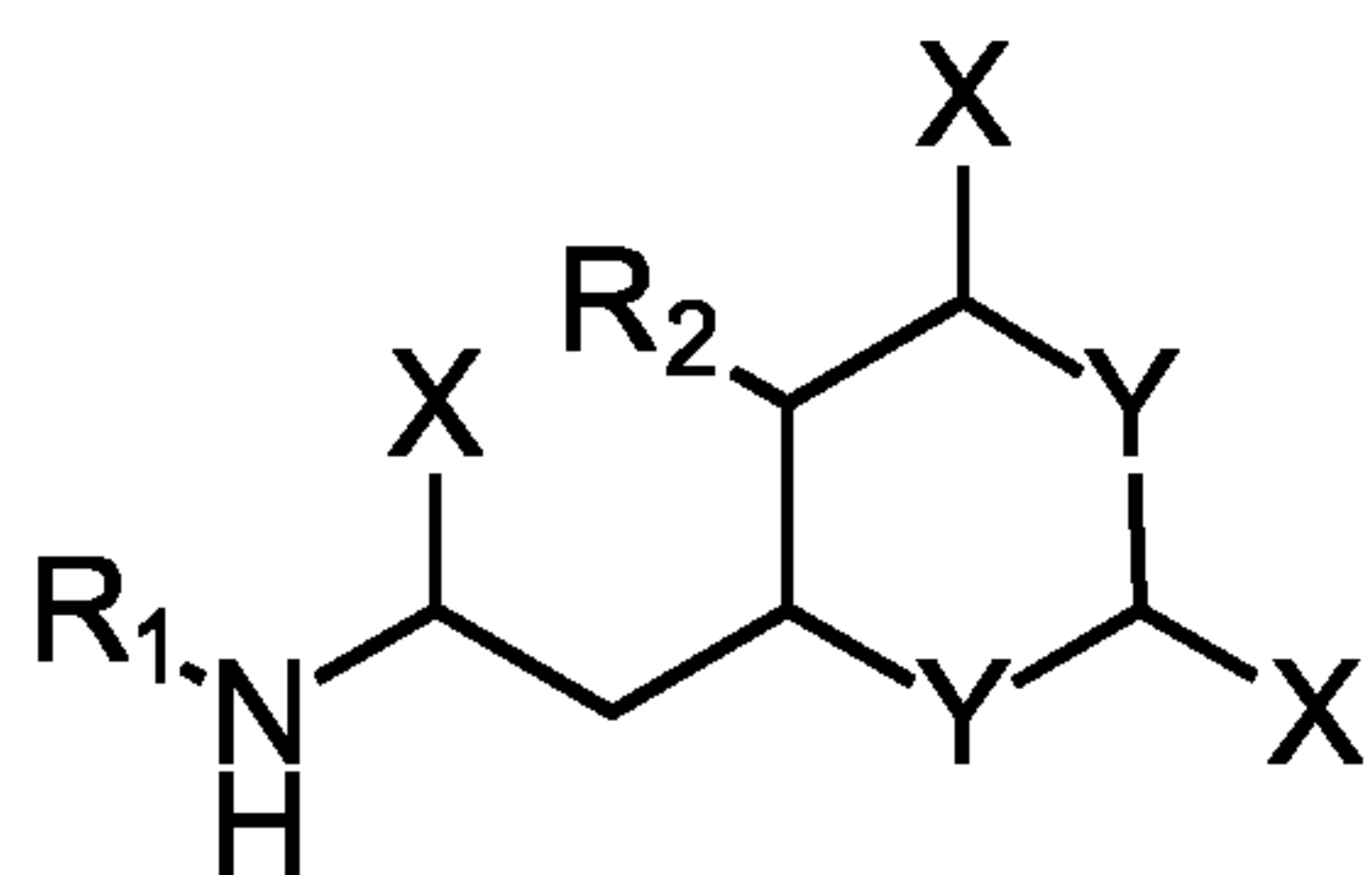
X represents a substituted or unsubstituted heteroatom selected from N and S; and

Y represents a linear n₃₋₅ alkyl, a linear n₃₋₅ alkenyl, or a linear n₃₋₅ alkynyl.

Compound 2 comprises for example the following modifications:



Compound 3 is represented by the following general Formula (III)



Formula (III)

wherein,

R₁ represents a hydrogen, linear n₃₋₅ alkyl, a linear n₃₋₅ alkenyl, a linear n₃₋₅ alkynyl, a linear n₃₋₅ hydroxylalkyl, a linear n₃₋₅ hydroxylalkenyl, a linear n₃₋₅ hydroxylalkynyl, a linear

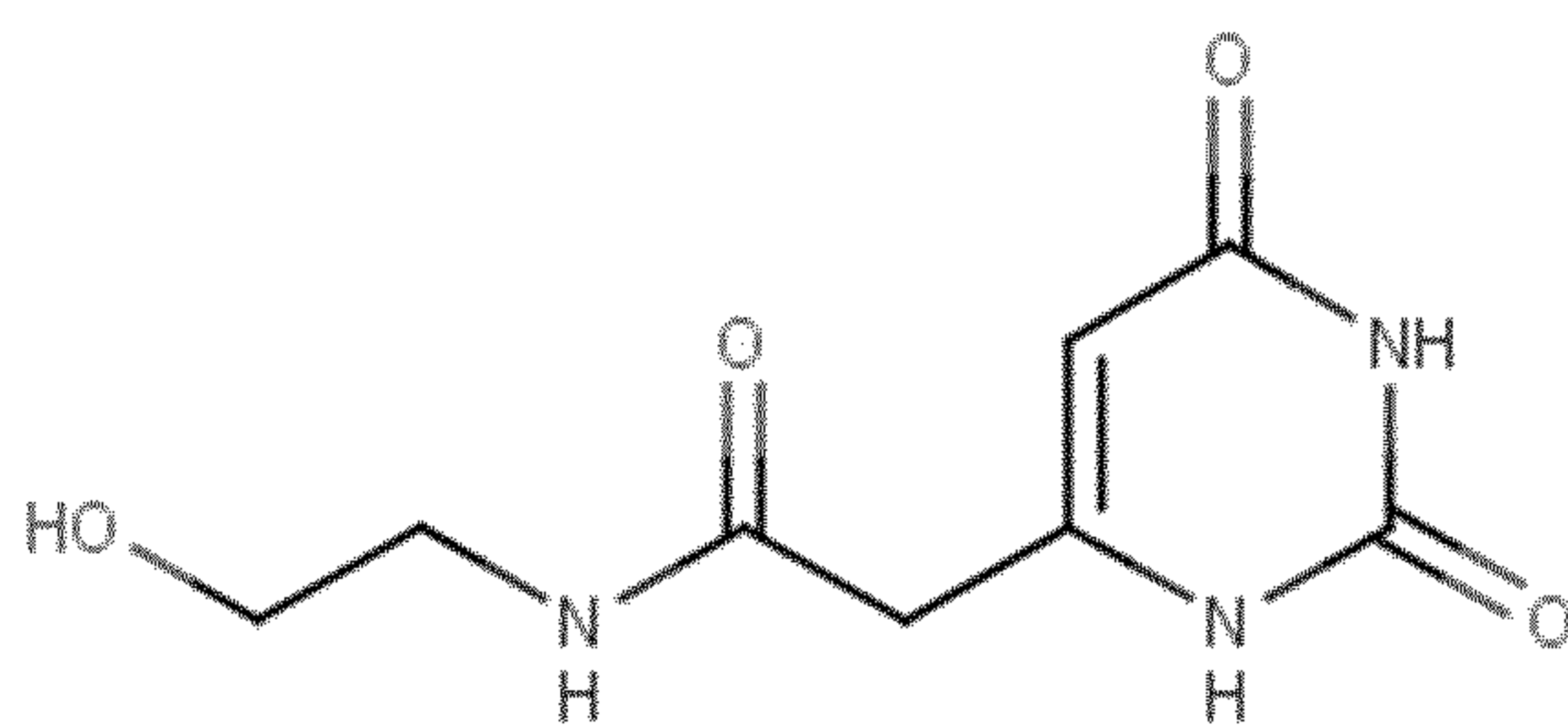
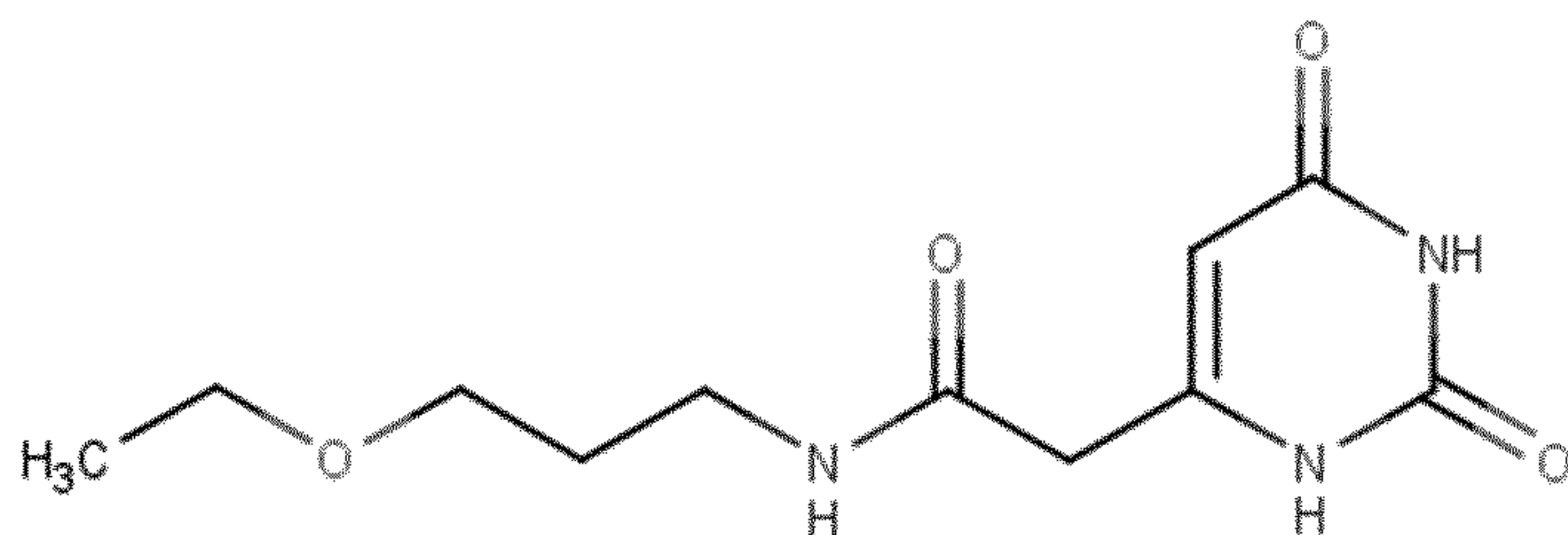
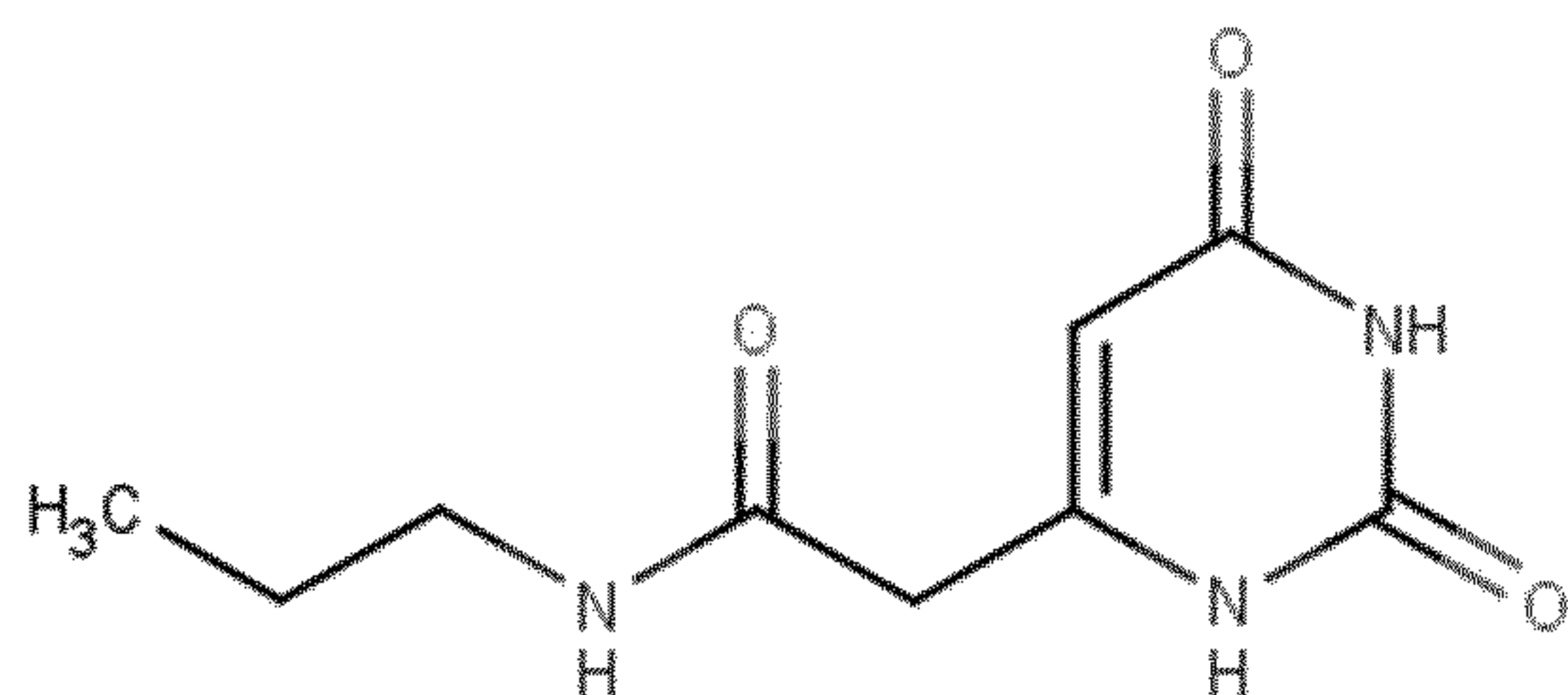
n₃₋₅ ethoxyalkyl, a linear n₃₋₅ ethoxyalkenyl, or a linear n₃₋₅ ethoxyalkynyl;

5 R₂ represents a hydrogen, halogen, alkyl, alkenyl, or alkynyl;

X represents a hydroxyl or a =O group;

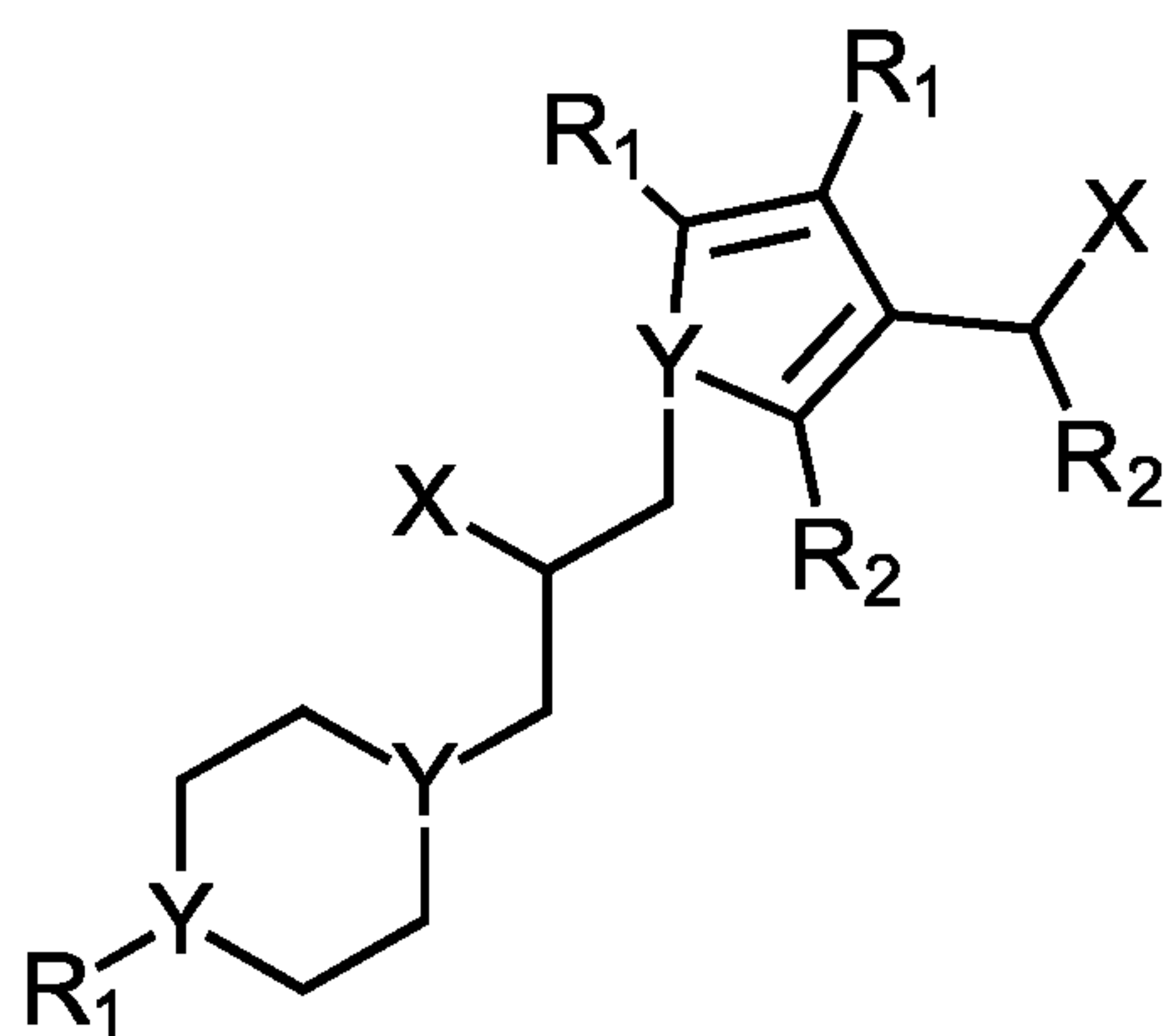
Y represents, independent of each other, a heteroatom selected from N and S, wherein the heteroatom may form a double bond with a neighboring carbon atom.

10 Compound 3 comprises for example the following modifications:



15

Compound 4 is represented by the following general Formula (IV)



Formula (IV)

wherein,

R₁ represents, independent from each other, a hydrogen, halogen, alkyl, alkenyl, or alkynyl;

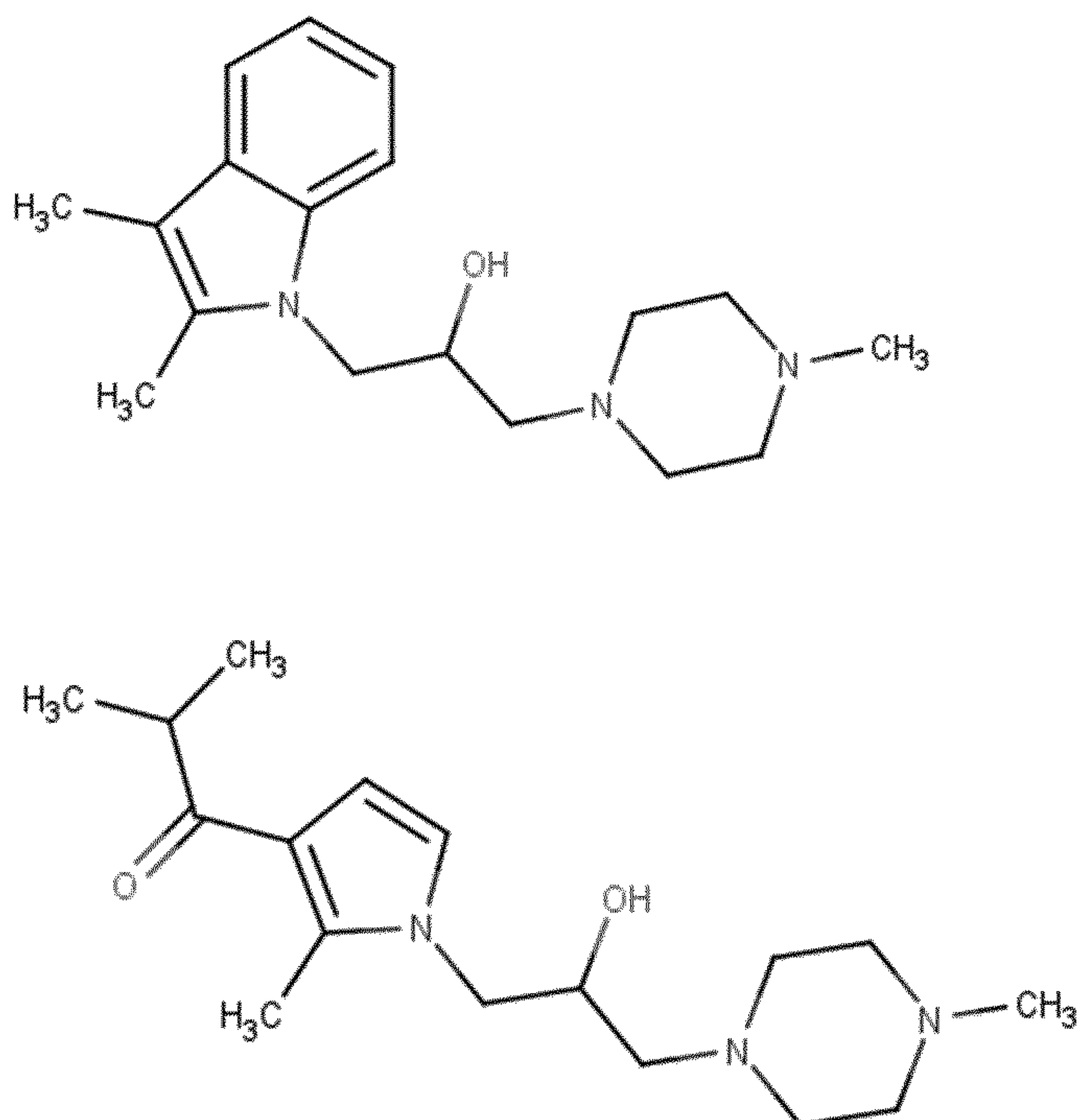
- 5 R₂ represents, independent from each other, a hydrogen, halogen, alkyl, alkenyl, alkynyl or wherein the R₂ residues together form a n₅₋₇ cycloalkyl or an n₅₋₇ aryl;

X represents a hydrogen, a hydroxyl or a =O group;

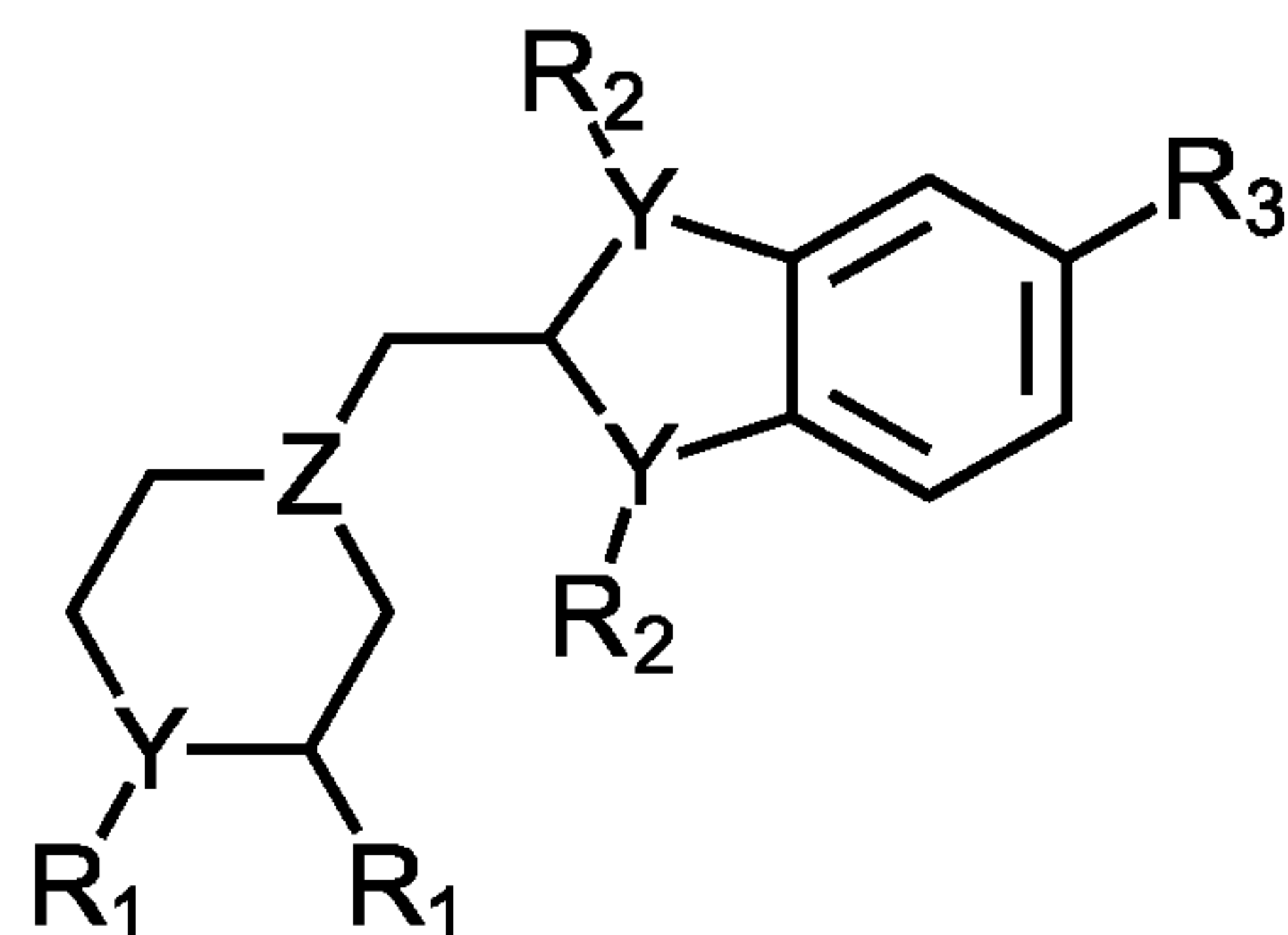
Y represents, independent from each other, a carbon atom or a heteroatom selected from N and S, wherein the heteroatom may form a double bond with a neighboring carbon

10 atom.

Compound 4 comprises for example the following modifications:



Compound 5 is represented by the following general Formula (V)



Formula (V)

5 wherein,

R₁ represents, independent of each other, a hydrogen, alkyl, alkenyl, or alkynyl, or wherein two R₁ form a cycloalkyl or an aryl;

R₂ represents, independent of each other, a hydrogen, alkyl, alkenyl, or alkynyl;

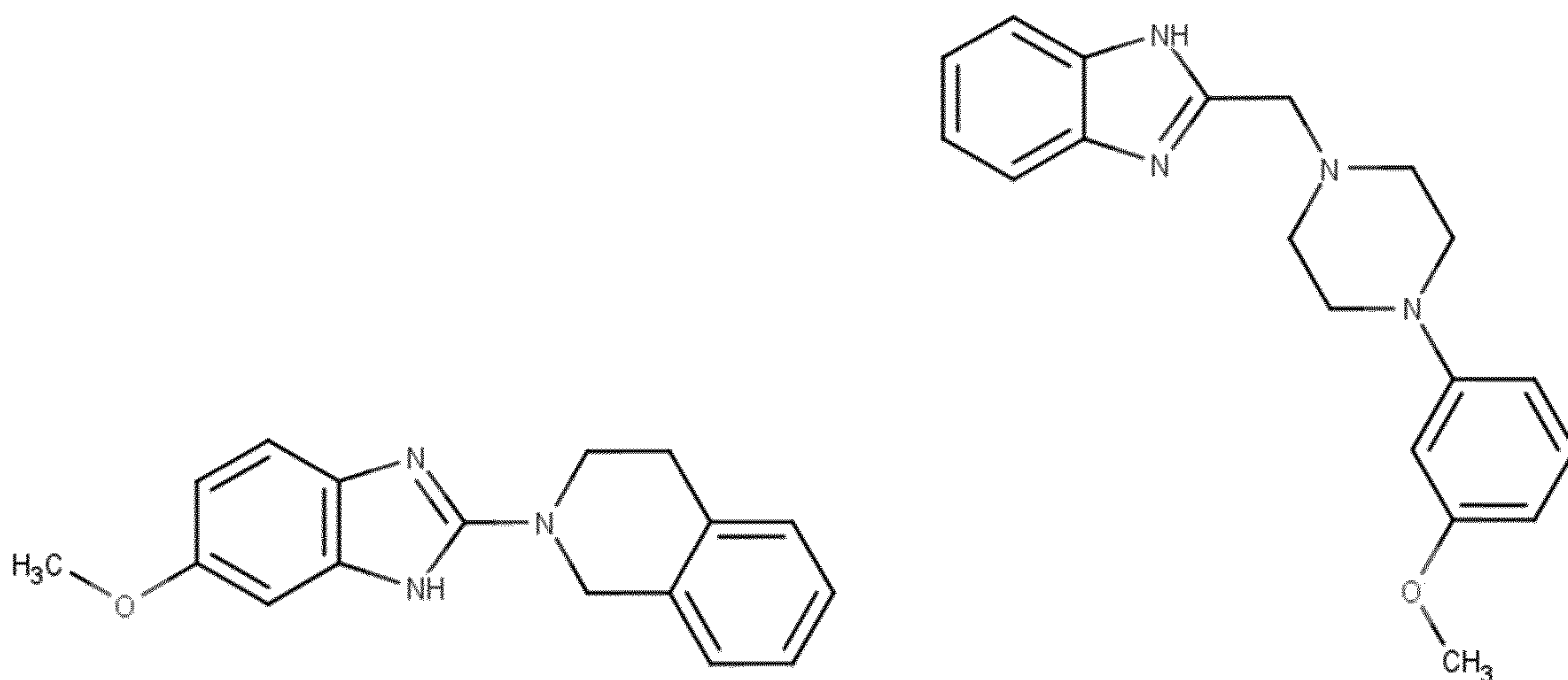
R₃ represents a hydrogen, halogen, ethoxyl, alkyl, alkenyl, or alkynyl;

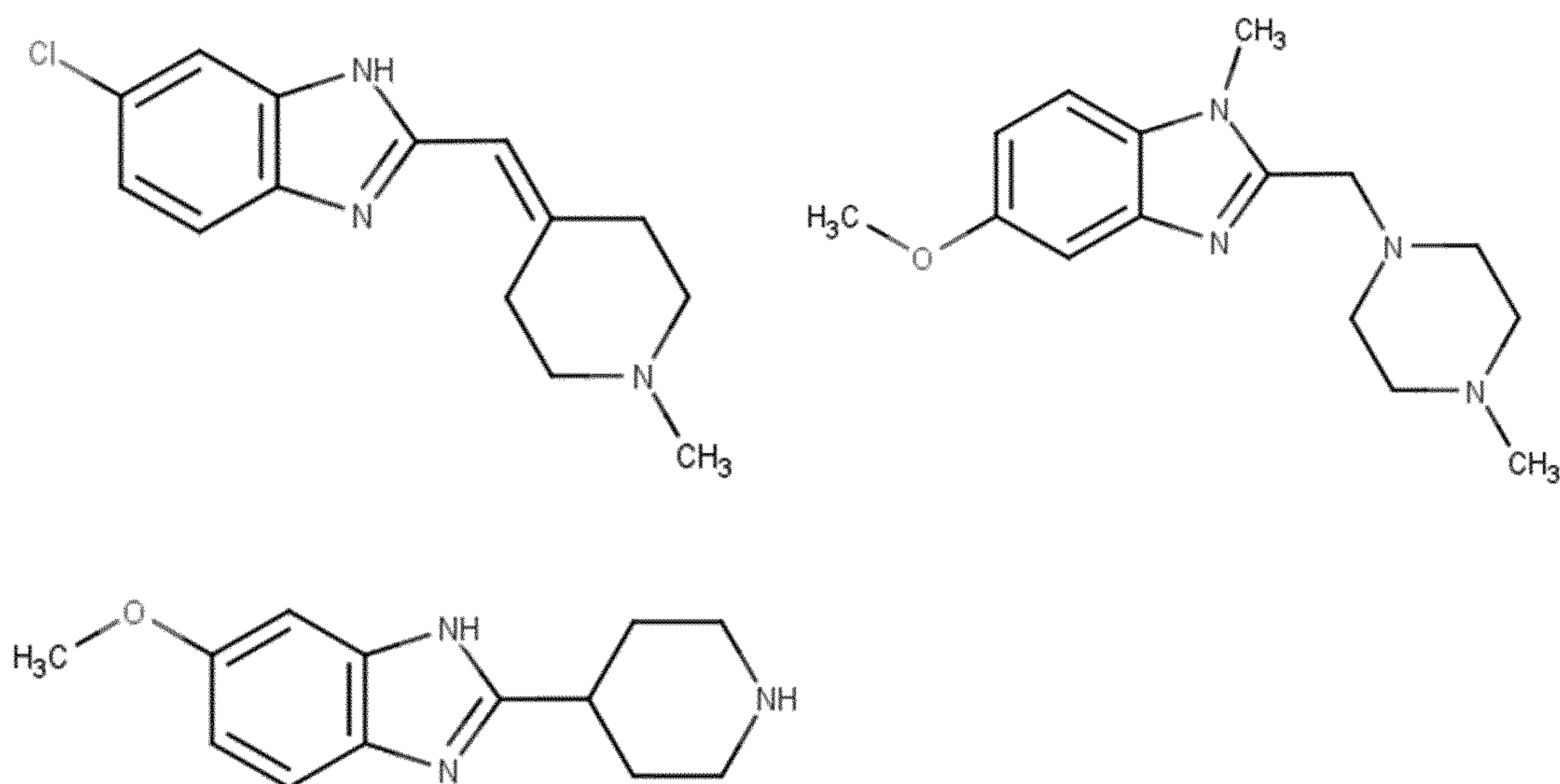
10 Y represents, independent of each other, a heteroatom selected from N and S, wherein the heteroatom may form a double bond with a neighboring carbon atom; and

Z represents a carbon atom or N.

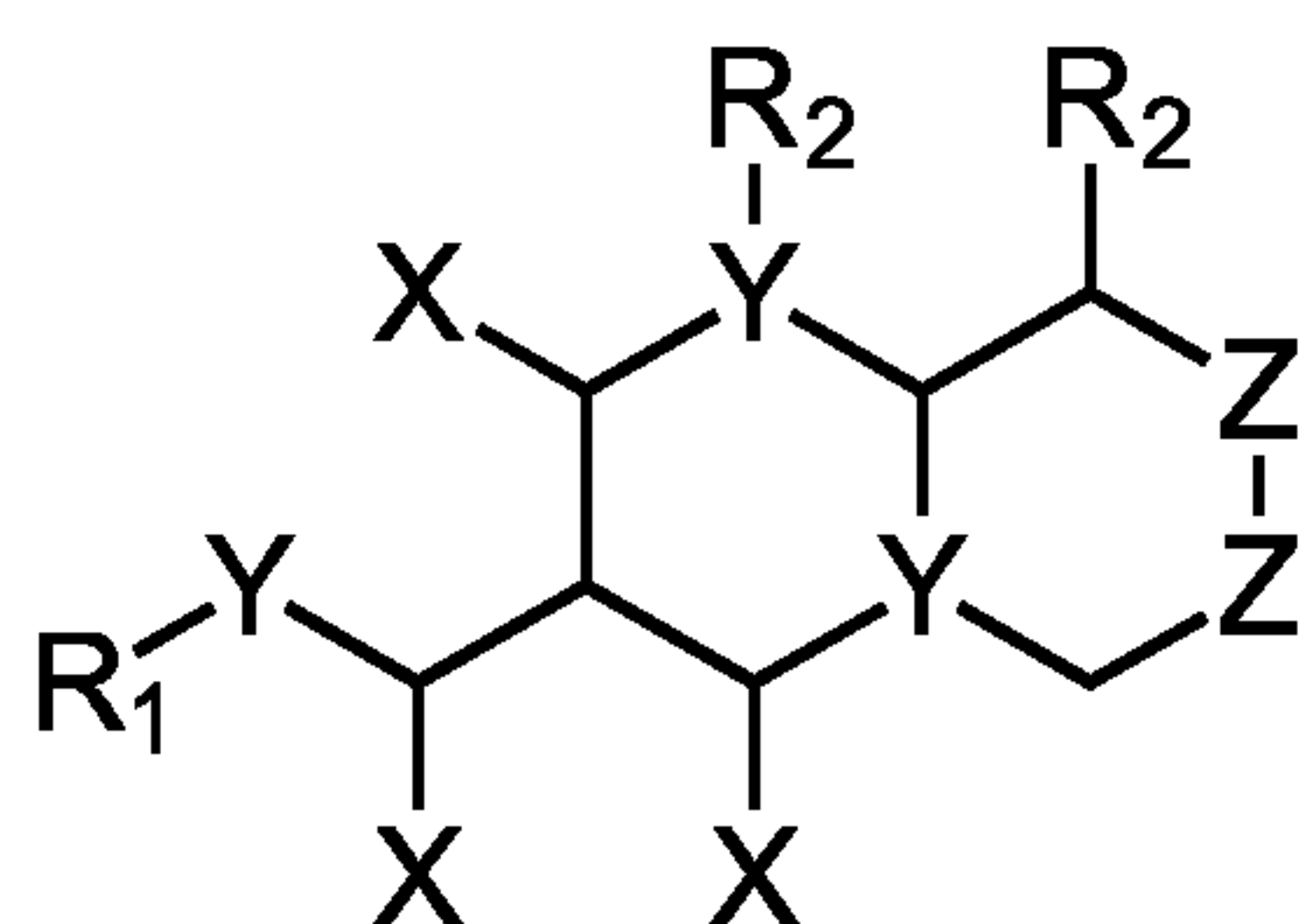
Compound 5 comprises for example the following modifications:

15





Compound 6 is represented by the following general Formula (VI)



Formula (VI)

wherein,

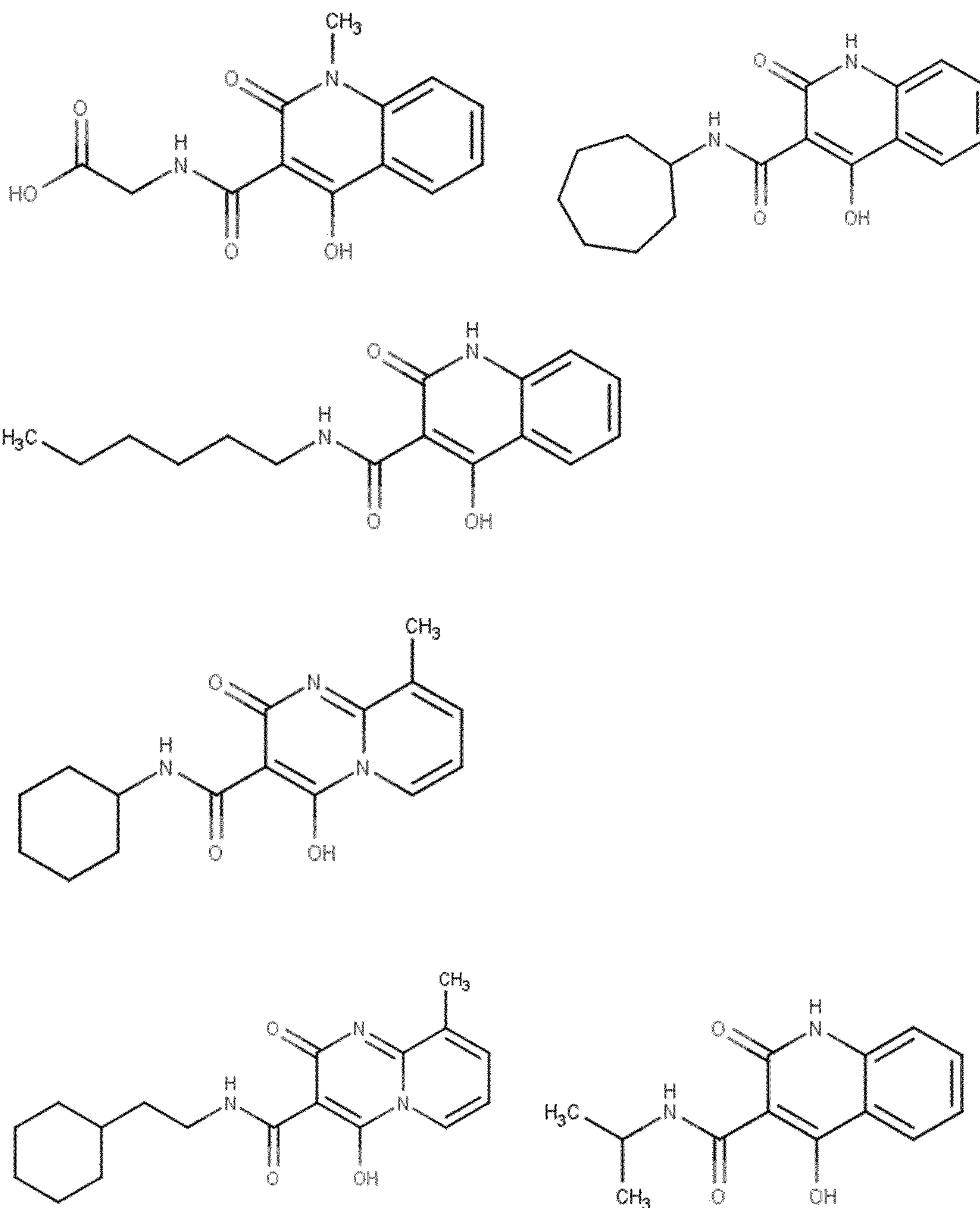
R_1 represents a linear n_{2-6} alkyl, a branched n_{3-6} alkyl, a linear n_{2-6} alkenyl, a branched n_{3-6} alkenyl, a linear n_{2-6} alkynyl, a branched n_{3-6} alkynyl, a n_{5-7} cycloalkyl, a n_{5-7} cycloalkenyl, a n_{5-7} cycloalkynyl, a n_{5-7} arylalkyl, a n_{1-3} alkylcarboxyl;

R_2 represents, independent of each other, a hydrogen, alkyl, alkenyl, or alkynyl;

Y represents, independent of each other, a heteroatom selected from N and S, wherein the heteroatom may form a double bond with a neighboring carbon atom; and

Z represents, independent of each other, a carbon atom which may form a double bond with a neighboring carbon atom.

Compound 6 comprises for example the following modifications:



5

The present invention is not only directed to the non-hydrophobic compounds of Table 1, to the tautomers and stereoisomers thereof, but also to chemically modified forms of these compounds as mentioned for example above.

- 10 The metastasis preventable and/or treatable by use of a compound of the present invention is based on any type of primary tumor for example melanoma, breast cancer, glioma, pancreas carcinoma, and colon carcinoma. In one embodiment, the primary

tumor is a late stage tumor. The metastasis is for example located in the liver, lung, bone, colon, stomach, nerves, lymph nodes, skin and/or brain.

The cartilage defect preventable and/or treatable by use of a compound of the present invention is for example a degenerative disorder of cartilage selected from rheumatoid arthritis, degeneration of cartilage in a joint, degenerative disc disease, meniscus tears, anterior crucial ligament (ACL) injury, arthritis, osteoarthritis, psoriatic arthritis, juvenile chronic arthritis, rhizomelic arthritis, rheumatoid poly-arthritis, synovitis and villonodular synovitis.

The dimerization of the MIA protein is for example measured by a heterogeneous transition metal-based fluorescence polarization (HTFP) assay, wherein the ratio P/P_0 is measured. P is the fluorescence polarization signal of a MIA protein labeled with a transition metal complex in the presence of substrate bound MIA-protein and of a compound to be tested. P_0 is the fluorescence polarization signal of free MIA-protein labeled with said luminescent transition metal complex in the absence of substrate bound MIA-protein and in the absence of the compound. In the absence of the compound, usually, the labeled MIA-protein would interact with the substrate bound MIA-protein, which, in turn, would contribute to a reduction in rotational mobility of the labeled MIA-protein, and therefore, the fluorescence polarization signal would increase upon such interaction. If, additionally, a compound is present that interferes with such interaction, no or little dimerization/aggregation occurs and no or little increase in fluorescence polarization signal would be detected. The smaller or even more negative P/P_0 is, the stronger such interference with dimer formation and aggregation is, and the better such compound prevents or breaks up dimerization/aggregation of MIA protein. In an alternative embodiment, binding of the compound to MIA protein is determined by NMR, e.g., heteronuclear NMR, e.g., ^{15}N - ^1H -HSQC-NMR.

In the present invention at least one compound of 1 to 270 of Table 1, a tautomer, a stereoisomer or a chemically modified compound thereof is administered to a subject, or in another embodiment two or more of these compounds, tautomers, stereoisomers and/or chemically modified compounds thereof are administered to a subject for use in preventing and/or treating of metastasis or a cartilage defect. The compounds are administered at the same time or consecutively.

In one embodiment the size of compounds of the present invention is increased by compound growing or compound linking. Regarding fragment growing an initial compound is steadily build up to explore favorable interactions with adjacent regions of the dimerization site; and regarding compound linking compounds
5 of the present invention (e.g., two or more for example 3, 4, 5, 6, 7, 8, 9, 10) are connected to each other, i.e., are coupled with or without a linker.

In an embodiment of the present invention the compounds are hydrophobic, in another embodiment the compounds are hydrophilic or neutral.

10

In another embodiment one or more compounds of the present invention are administered to a subject in combination with a chemotherapeutic which is any chemical agent, e.g., naturally occurring or synthesized, effective in the treatment of cancer such as Vermurafenib, Ipilimumab, Trametinib, Dabradenib, Dacarbazine, Paclitaxel,
15 Carboplatin, Interferon-alpha and Aldesleukin, TGF alpha, TGF beta, interleukin etc.. The compound and the chemotherapeutic are administered at the same time or consecutively.

20

The present invention further refers to a pharmaceutical composition for use in preventing and/or treating of metastasis caused by the dimerization of melanoma inhibitory activity (MIA) protein and/or a cartilage defect, wherein regeneration is inhibited by MIA dimerization, wherein the pharmaceutical composition comprises at least one compound selected from the group consisting of compound 1 to 270 of Table 1, tautomers, stereoisomers and/or chemically modified compounds thereof, and a
25 pharmaceutically acceptable carrier and/or solvent. The pharmaceutical composition may further comprise a chemotherapeutic or any substance, including cells such as chondrocytes, positively influencing cartilage growth and regeneration, respectively.

Examples

30

The following examples show the present invention in more detail, however, the invention is not limited to these examples.

Example 1

35

Fragment-based *in silico* screening was applied to identify the small, fragment-sized compounds of the present invention that inhibit MIA. The suggested structures were

screened *in vitro* and modular synthesis strategies were developed for the most promising molecules.

NMR titration experiment

- 5 Derived from HSQC titrations of ¹⁵N-labeled MIA with AR71 (a peptide having the amino acid sequence FHWRYPLPLPGQ), the amino acids CYS17, SER18, TYR47, GLY66, ASP67, LEU76, TRP102, ASP103, and CYS106 were found to exhibit strong shift perturbations (Schmidt, J., A. Riechers, and A.K. Bosserhoff, *MIA - a new target protein for malignant melanoma therapy*. *Histol Histopathol*, 2013. **28**(4): p. 421-6) and
- 10 were defined as interacting residues in an *in silico* protein-peptide docking of MIA and AR71. The resulting model shows the binding of the peptide in the hydrophobic cleft, which forms part of the dimerization domain.

Virtual screening

- 15 The 1.4 Å resolution crystal structure of human MIA protein 1I1J (Lougheed, J.C., et al., *Structure of melanoma inhibitory activity protein, a member of a recently identified family of secreted proteins*. *Proceedings of the National Academy of Sciences*, 2001, **98**(10): p. 5515-5520) and the first model of each of the NMR solution structures 1K0X (Lougheed, J., P. Dommelle, and T. Handel, *Solution structure and dynamics of melanoma*
- 20 *inhibitory activity protein*. *Journal of Biomolecular NMR*, 2002, **22**(3): p. 211-223) and 1HJD (Stoll, R., et al., *The extracellular human melanoma inhibitory activity (MIA) protein adopts an SH3 domain-like fold*. 2001. **20**(3): p. 340-349) were used as distinct MIA protein receptor conformations for the virtual screening experiment. The same amino acid residues that were used in the protein-peptide docking were provided as
- 25 input to the molecular docking software in order to guide the automated detection and definition of putative binding sites in the MIA protein.

In each of the three MIA protein conformations the same single putative binding site was detected. The site is located in a cleft near the distal loop (residues 69 to 75) and is

30 framed by the solvent exposed “disulfide loop” (residues 13 to 19), the turn of the RT loop (residues 35-38), and the C-terminal residues 102 to 106. Most of the amino acid residues identified to interact with the AR71 peptide by induced chemical shift changes constitute to the putative binding site, with the exception of TYR48, GLY66, and ASP67, which are located on the distal loop and thus too far away. Differences in the arrangement of the

structural elements surrounding the cleft within each of the three distinct MIA receptor conformations result in a different volume and shape of the putative binding site.

Fragment library preparation commenced with 28751 fragment structures that were contained within the unprocessed fragment library catalogue obtained from the compound provider. Filtering for salts, unwanted chemistry and functional groups, as well as duplicate structures led to the removal of 1, 2577 and 127 structures, respectively. Each of the remaining 26046 structures was subjected to complete enumeration of its protonation and tautomeric states, which led to 87270 protomer variants for all fragment structures. Subsequently, all stereoisomers were enumerated for the protomer variants, which resulted in 124590 fragment variants. Generation of 3D conformations allowed for up to five alternate ring conformations per fragment variant and produced a final library of 217176 fragment variant conformations, which were docked into the three MIA receptor conformations with their respective putative binding sites.

The resulting docking poses were subjected to a filtering and ranking workflow, which, for each MIA protein conformation, determined a list of fragments ranked by the empirical scoring function of the docking software. The ten highest-ranking fragments for each MIA receptor conformation were selected for experimental testing.

All listed compounds that were screened for their ability to interfere with the MIA-MIA interaction (Table 1) in a heterogeneous transition metal-based fluorescence polarization (HTFP) assay previously developed (Riechers, A., et al., *Heterogeneous transition metal-based fluorescence polarization (HTFP) assay for probing protein interactions*. Biotechniques, 2009. 47(4): p. 837-44). This screening revealed significant decreases in the fluorescence polarization of Ru-(bpy)₃, labelled MIA for compounds 1, 2, 3, 4, 5, and 6; the results are shown in Fig. 3. The molecular structures of the compounds showing interference with the MIA-MIA interaction in the HTFP assay are shown in Table 1.

Example 2

Boyden chamber migration assays (Stoll, R., Lodermeier, S. & Bosserhoff, A. K. Detailed analysis of MIA protein by mutagenesis. Biol Chem 387, 1601 - 1606, (2006)) using human Mel-Im melanoma cells demonstrated a reduction in the MIA activity on melanoma cell migration in the presence of compounds 1, 2, 3, 4, 5, and 6 at a concentration of 1 μ M as shown in Fig. 4.

The melanoma cell line Mel-Im, established from a human metastatic bioptic sample (generous gift from Dr. Johnson, University of Munich, Germany) was used for the Boyden chamber migration experiments. All cells were maintained in DMEM (PAA, Pasching, Germany) supplemented with penicillin (400 U/mL), streptomycin (50 µg/mL), L-glutamine (300 µg/mL) and 10% fetal calf serum (Pan Biotech GmbH, Aidenbach, Germany) and split in a 1:6 ratio every three days. Migration assays were performed in Boyden Chambers containing polycarbonate filters with 8-µm pore size (Neuro Probe, Gaithersburg, MD, USA) essentially as described. MIA was added to the cell suspension at a final concentration of 200 ng/mL. Selected compounds were used at a final concentration of 1 µM. Experiments were carried out in triplicates and repeated at least three times.

Example 3

In order to assess whether compounds of the present invention show any adverse effects on normal cells, human fibroblasts and kidney cells were treated with the compounds 1, 2, 3, 4, 5, and 6, respectively, *in vitro* at a concentration of 7.8 µM in analogy to previous studies (Schmidt, J., et al., *Targeting melanoma metastasis and immunosuppression with a new mode of melanoma inhibitory activity (MIA) protein inhibition*. PLoS One, 2012. 7(5): p. e37941; Riechers, A., et al., *Heterogeneous transition metal-based fluorescence polarization (HTFP) assay for probing protein interactions*. Biotechniques, 2009. 47(4): p. 837-44). There were no negative effects on the proliferation of human fibroblasts (Fig. 5A) and kidney cells (Fig. 5C) as well as cellular attachment of the same cell types (Fig. 5B and 5D) in the presence of the compounds.

A significant decrease in proliferation of the human melanoma cell line Mel Im after treatment with compound 1 was observed as presented in (Fig. 6). A phenomenon that could not be found with the other compounds 2, 3, 4, 5 or 6.

Claims

- 1) Non-hydrophobic compound for use in preventing and/or treating of metastasis caused by the dimerization of melanoma inhibitory activity (MIA) protein or of a cartilage defect,
5 wherein regeneration is inhibited by MIA dimerization wherein the compound is selected from the group consisting of compound 1 to 300 of Table 1, tautomers, stereoisomers and chemically modified compounds thereof.
- 2) Compound for use according to claim 1, wherein the compound is compound 1, 2, 3, 4,
10 5 and/or 6.
- 3) Compound for use according to claim 1 or 2, wherein the metastasis is based on a primary tumor selected from the group of MIA-expressing tumors.
- 15 4) Compound for use according to claim 3, wherein the metastasis is based on a primary tumor selected from the group consisting of melanoma, breast cancer, glioma, pancreas carcinoma, and colon carcinoma.
- 5) Compound for use according to any one of claims 1 to 4, wherein the metastasis is
20 located in the liver, lung, bone, colon, stomach, nerves, lymph nodes, skin and/or brain.
- 6) Compound for use according to any one of claims 1 to 5, wherein the compound is administered orally.
- 25 7) Compound for use according to any one of claims 1 to 6 in combination with a chemotherapeutic.
- 8) Compound for use according to claim 7, wherein the chemotherapeutic is selected from the group consisting of Vermurafenib, Ipilimumab, Trametinib, Dabradenib,
30 Dacarbazine, Paclitaxel, Carboplatin, Interferon-alpha and Aldesleukin.
- 9) Compound for use according to claim 7 or 8, wherein the compound and the chemotherapeutic are administered at the same time or consecutively.

10) Pharmaceutical composition for use in preventing and/or treating of metastasis caused by the dimerization of melanoma inhibitory activity (MIA) protein and/or a cartilage defect, wherein regeneration is inhibited by MIA dimerization, wherein the pharmaceutical composition comprises at least one compound selected from the group
5 consisting of compound 1 to 300 of Table 1, tautomers and stereoisomers thereof, and a pharmaceutically acceptable carrier and/or solvent.

11) Pharmaceutical composition for use according to claim 10, further comprising a chemotherapeutic.

Fig. 1: Model of MIA dimer forming dimerization (binding) site

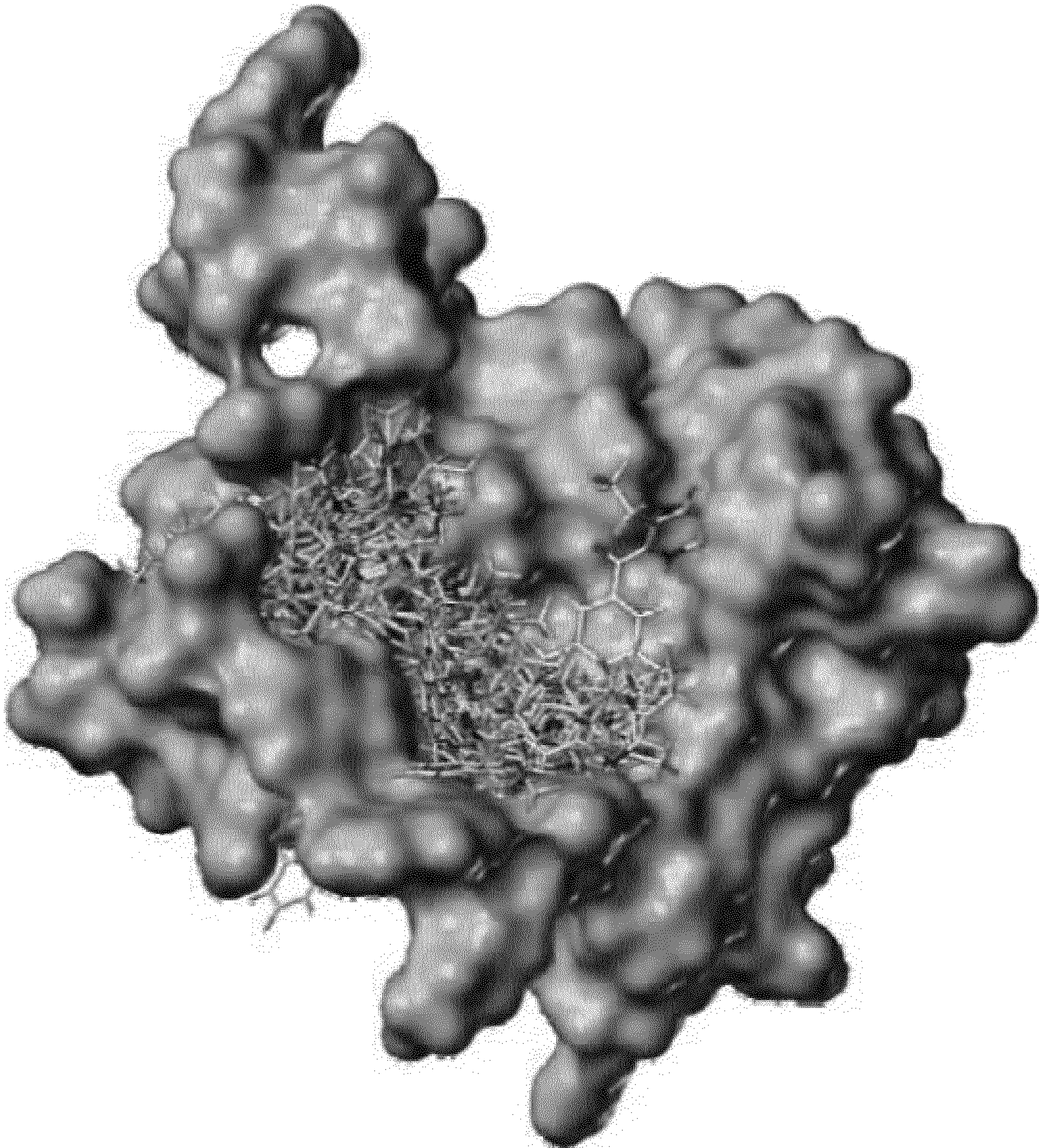


Fig. 2A: Predicted passing of the blood-brain-barrier of compound 3

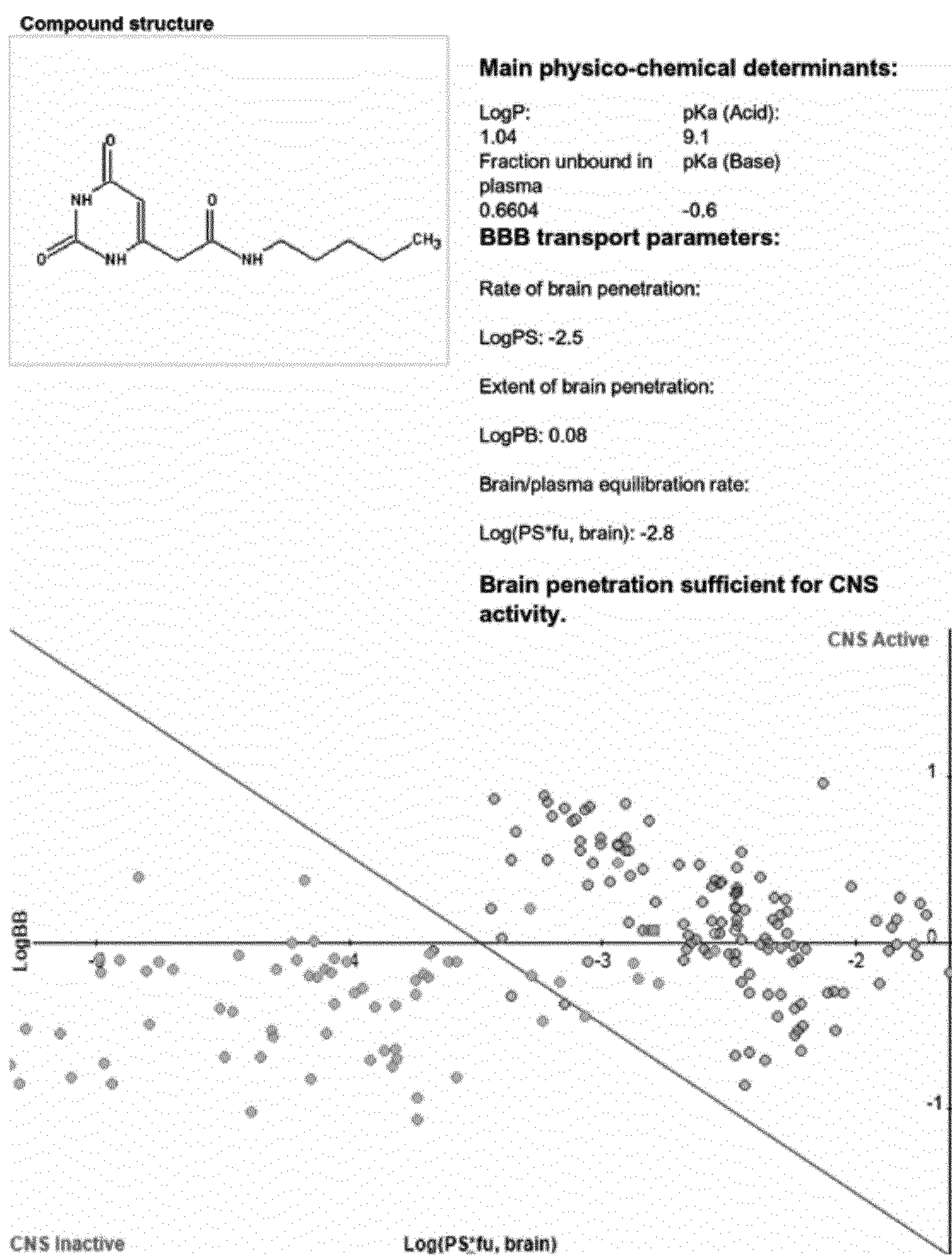


Fig. 2B: Predicted oral availability of compound 3

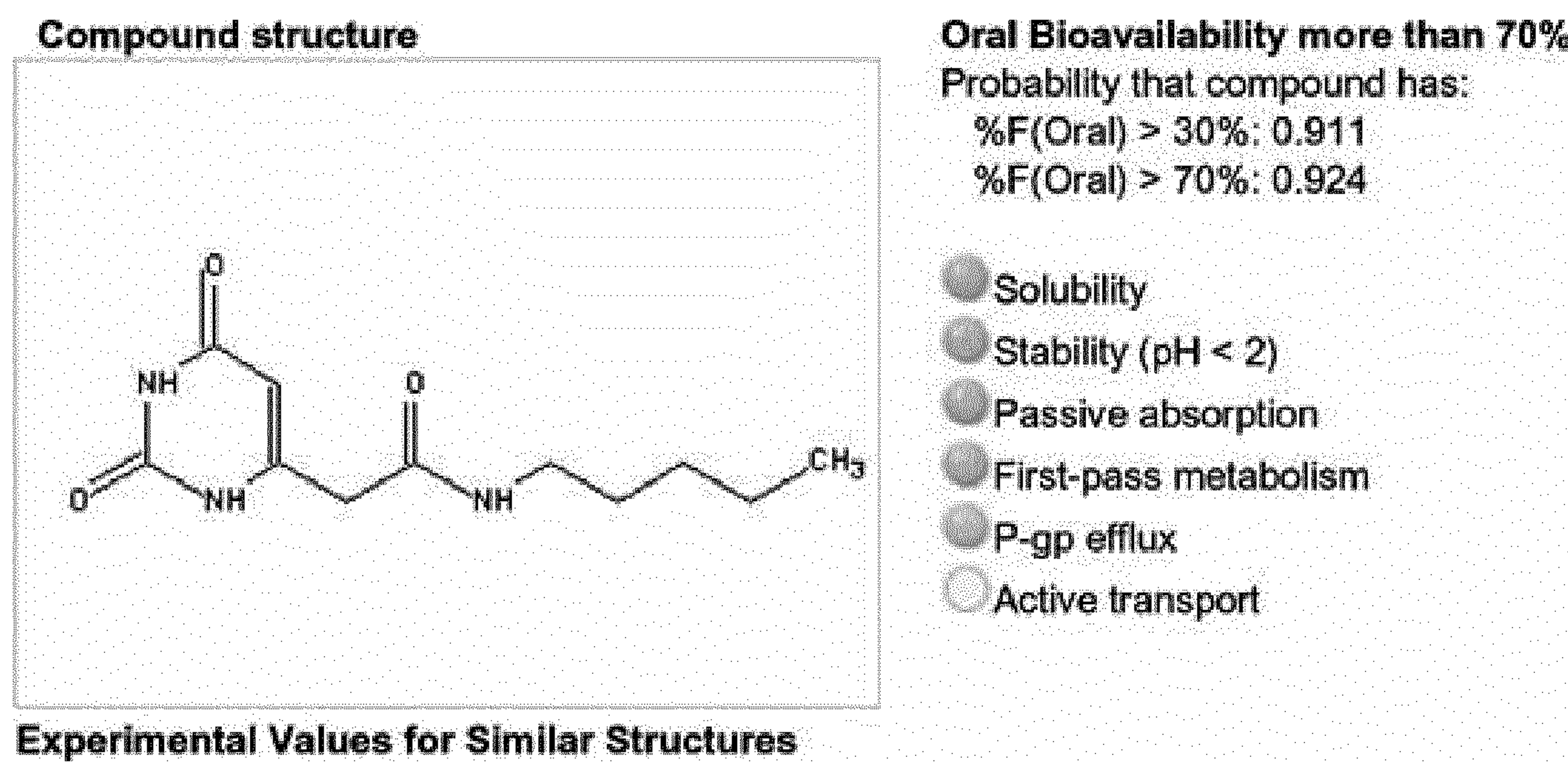


Fig. 2C: Predicted systemic toxicity of compound 3

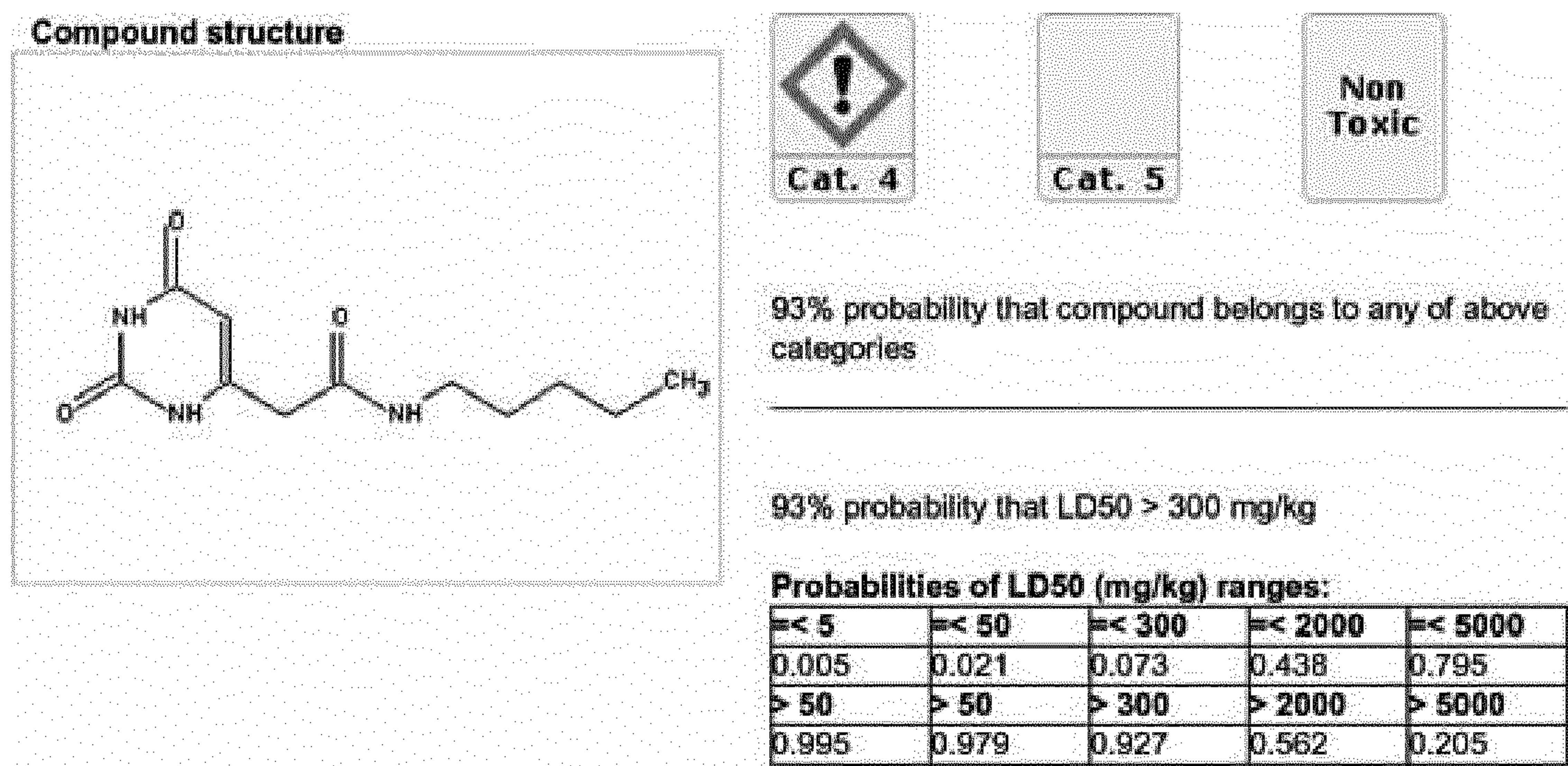


Fig. 3: Inhibition of MIA dimerization measured in HTFP assay

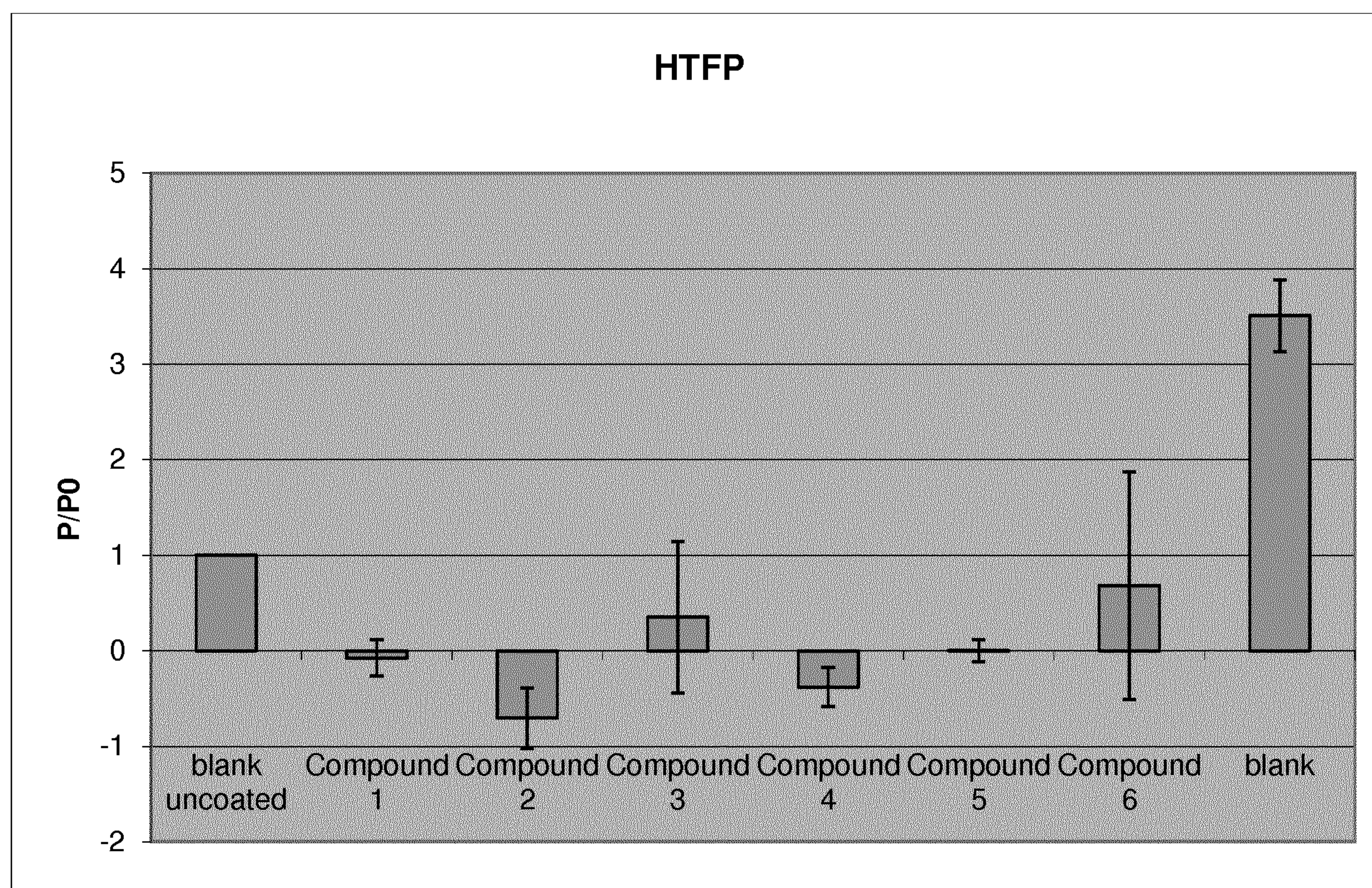


Fig. 4: Inhibition of MIA activity

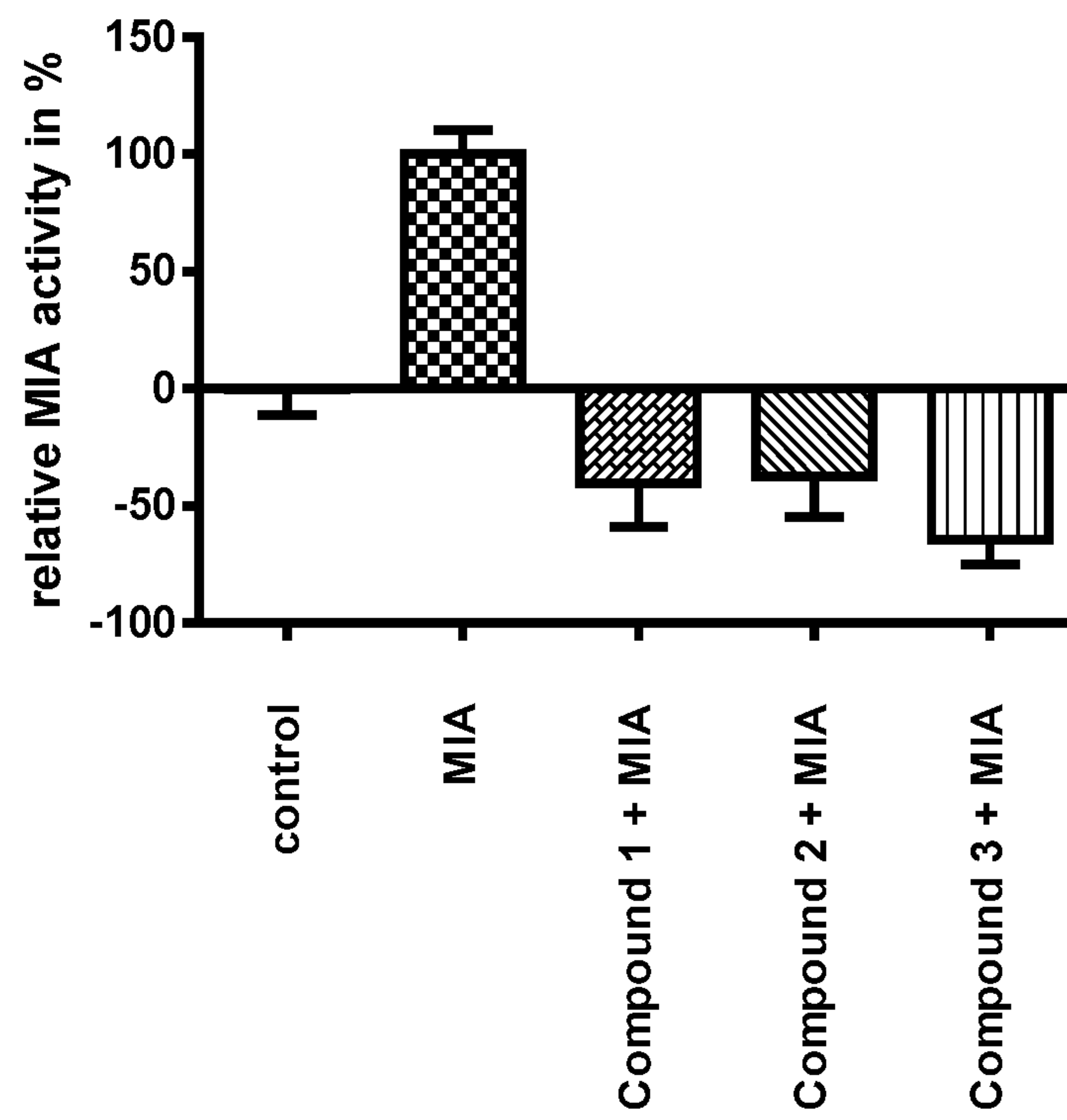


Fig. 5A and 5B: Effect of compounds 1 to 6 on fibroblasts

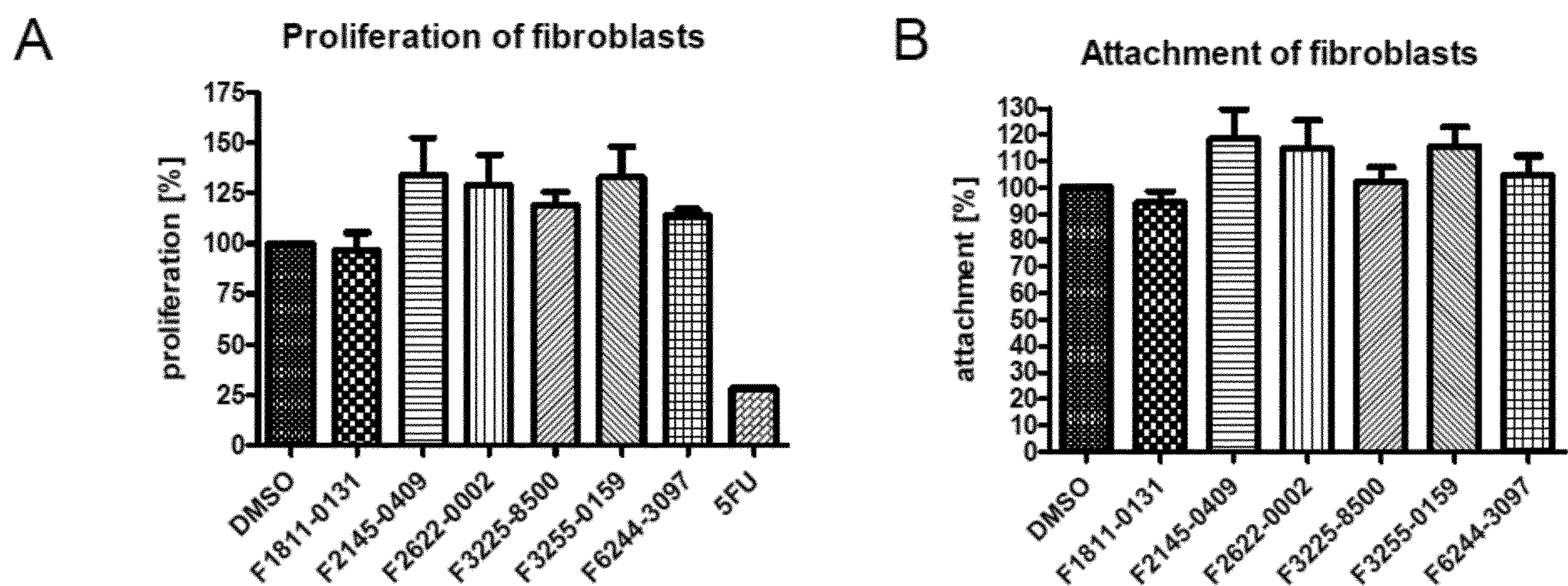


Fig. 5C and 5D: Effect of compounds 1 to 6 on kidney cells

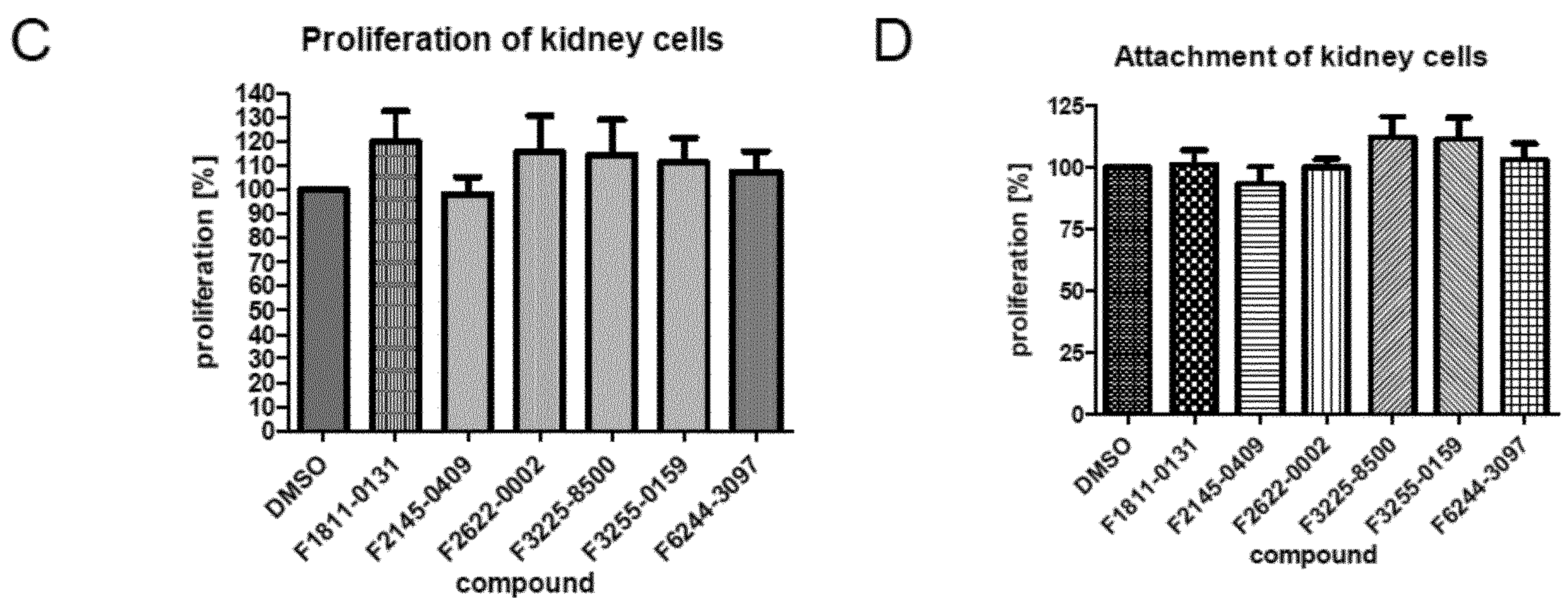


Fig. 6: Effect of compound 1 on proliferation of melanoma cells

