

Conformation–Function Relationship of Vitamin D: Conformational Analysis Predicts Potential Side-Chain Structure

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In previous studies, we have grouped regions in space occupied by the vitamin D side chain into four: A, G, EA, and EG. We showed that the receptor (VDR) affinity of $1\alpha,25$ -dihydroxyvitamin D₃ derivatives increases, in terms of side-chain region, in the order EG, G, A, and EA. We called this the active space group concept. In the present study, we used this active space group concept to analyze the conformation–activity relationship of about 40 representative potent $1\alpha,25$ -dihydroxyvitamin D₃ analogues. We initially listed structural modifications in the side chain of potent vitamin D analogues and estimated their potency factor. Possible side-chain conformations of representative analogues were calculated by the molecular mechanics method and plotted on a dot map compared with the regions A, G, EA, and EG. The cell-differentiating potency of the analogues was correlated with our active space group concept with few exceptions. Among potent analogues with a natural configuration at C(20), the side chains of those with a 22-oxa, 22-ene, 16-ene, or a 18-nor modification were located in front of region EA (termed F). The side chains of the most potent 20-epi-22-oxa-24-homovitamin D analogues were concentrated at the left side of the EA region (L-EA). Thus, the side chains of almost all potent analogues were distributed around the EA region, and potency increased in the order A, F, EA, and L-EA.

Introduction

Binding of a ligand to the receptor is the initial and most crucial step in activating gene transcription mediated by a ligand-dependent nuclear transcription factor. How molecular recognition occurs at the interface of a ligand and the receptor is of interest to chemists. Control of this process is also key to developing potent and selective clinical agents.

Vitamin D exerts its function by binding to its nuclear receptor (VDR),¹ which is a member of the nuclear receptor superfamily.² In binding to VDR, the three hydroxyl groups of $1\alpha,25$ -dihydroxyvitamin D₃ ($1,25$ -(OH)₂D₃, **1**) play important roles. In particular, those of the 1α - and 25-hydroxyl groups are critical, as their removal diminishes the VDR affinity by 1/500–1/1000.³

Over 300 active vitamin D analogues have been synthesized⁴ since the discovery of $1,25$ -(OH)₂D₃ (**1**),⁵ and some are in clinical use to treat metabolic bone diseases and skin disorders such as psoriasis.⁶ Nearly 100 analogues with higher activities than that of the parent natural ligand (**1**) have been identified, and most of them are modified around the side chain.⁴ In our initial structure–function studies of vitamin D,⁷ we focused our attention on the side chain. Vitamin D side chains are highly flexible as they possess the full carbon structure of cholesterol. Therefore the 25-hydroxyl group, which apparently plays a crucial role in anchoring vitamin D to the VDR, can occupy wide regions of space. The 20-epimer **2** of $1,25$ -(OH)₂D₃ (**1**) is much more potent than the natural vitamin,^{8a} although the side chain of these epimers appears to occupy different

regions of space. Okamura and Midland⁹ analyzed the conformational mobility of the side chain of these epimeric vitamins (**1** and **2**) and relatives by plotting the 25-oxygen position of possible conformations on a dot map. In these studies, they discriminated the contribution of the conformers using the energy window concept. To determine the VDR-bound side-chain conformation, we initially analyzed possible side-chain conformations of $1,25$ -(OH)₂D₃ (**1**) and its 20-epimer **2** by molecular mechanics and grouped the regions occupied by their 25-oxygen into four: A, G, EA, and EG (Figure 1).^{7a,b} We then designed and synthesized analogues in which the side-chain mobility was restricted in one of these regions, the diastereomers **3**–**6** at C(20) and C(22) of 22-Me- $1,25$ -(OH)₂D₃.^{7a,b} The VDR affinity of these conformationally restricted analogues increased in the order **6**, **3**, **4**, and **5** (1/100, 1/60, 1/3, and 20, respectively, relative to $1,25$ -(OH)₂D₃, **1**). In terms of the region, the potency increased in the order EG, G, A, and EA.^{7a} From these results, we postulated that $1,25$ -(OH)₂D₃ (**1**) binds to VDR when its side chain is located in the A region¹⁰ and that 20-epi- $1,25$ -(OH)₂D₃ (**2**) binds to VDR when its side chain is located in the EA region.

We next studied whether our active space group concept can predict the potency of known vitamin D analogues. In this study, we (1) listed structural modifications in the side chain (including C(17)) of vitamin D that elevate potency, (2) calculated the possible side-chain conformations of these analogues that were displayed on dot maps, and (3) investigated the relationship between the side-chain region and activity. The results not only support our idea of active areas of the side chain but also demonstrate that

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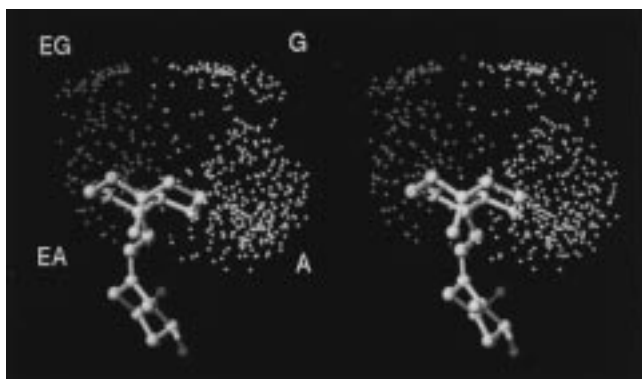


Figure 1. Grouping of the side-chain regions of 1,25-(OH)₂D₃ (**1**) and its 20-epimer (**2**) (stereoview). The mobile areas of the side chain of **1** (yellow) and **2** (cyan) are shown in a dot map and are grouped into regions: A and G regions in **1** and EA and EG regions in **2** (see ref 7a).

conformational analysis can predict the potency of vitamin D analogues.

Methods

Potent Vitamin D Analogues and Activation Factor.

We compared the potency of vitamin D derivatives in terms of their cell-differentiating activity (throughout this paper, potency indicates cell-differentiating activity unless otherwise noted). Affinity for VDR may be more appropriate for predicting receptor-bound conformation, but IC₅₀ values determined by competitive binding assays are not always reproducible for compounds having higher affinity than 1,25-(OH)₂D₃ (**1**). This is because attaining equilibrium among the receptor and two such ligands (the sample and radioactive standard) is difficult under assay conditions.¹¹

Table 1 lists structural modifications that enhance potency, together with representative compounds and activation factors estimated on the basis of cell-differentiation activity. For

simplicity, the analogues were restricted to those derived from 1,25-(OH)₂D₃ (**1**). The modifications that increase potency are (1) epimerization at the 20-position,⁸ (2) replacement of C(22)-H₂ by oxygen,¹² (3) introduction of double or triple bond(s) to the side chain¹³ or the 16-position,¹⁴ (4) removal of the angular C(18) methyl group,¹⁵ (5) elongation of the side chain,¹⁶ (6) methylation of the terminal C(26) and -(27) methyl groups,¹⁶ (7) perfluorination of C(24) methylene¹⁷ or the terminal gem-dimethyl groups,¹⁸ and (8) methylation at C(20).¹⁹ Combinations of these modifications enormously enhance potency as the compounds in Table 2 show. In most cases, the potency of compounds with multimodifications can be estimated by multiplying by the activation factor shown in Table 1.

Conformational Analysis. We analyzed the relationship between the potency of the analogues and the areas occupied by the 25-oxygen. (1) The possible side-chain conformations of each analogue were searched using the molecular mechanics method. (2) These possible conformations were plotted by positioning the 25-oxygen on a dot map where one dot corresponds to one-side chain conformation. (3) The dot map of each analogue was overlaid with those of 1,25-(OH)₂D₃ (**1**) and 20-epi-1,25(OH)₂D₃ (**2**), and then we analyzed the relationship between the side-chain area and cell-differentiating activity.

The conformations were analyzed using the software SYBYL (version 6.3, Tripos)²⁰ as reported.^{7a,b} Briefly, (1) analogue structures were constructed by modifying the crystallographic vitamin D₃ structure in the SYBYL database; then the global minimum-energy conformation of each analogue was generated by repeating rotations (360° at 60° intervals) of rotatable C–C and C–O bonds in the side chain (grid search in SYBYL) followed by optimization. (2) Using this minimum-energy conformation as the initial structure, all possible side-chain conformations were searched by rotating (360° at 30° intervals) the bonds in the side chain (indicated by numbers in the structures in Chart 1) (systematic search in SYBYL). In the systematic search, the van der Waals bump coefficient was set at 0.95 to eliminate infeasible conformations suffered from van der Waals repulsion. (3) All possible conformations were

Table 1. Side-Chain Modifications That Elevate Vitamin D Potency

| modification | abbrev | activation factor | representative compound | |
|-----------------------------------|---------------------------------------|-------------------|---|--|
| 20-epimerization | EP | 20–30 | 20-epi-1,25-(OH) ₂ D ₃ (2) ^{8a} | |
| 22-oxygen | 22O | 5–10 | 22-oxa-1,25-(OH) ₂ D ₃ (7) ¹² | |
| unsaturation [yne (Y) or ene (E)] | 22E | 1–2 | 22-ene-1,25-(OH) ₂ D ₃ (8) ^{13a} | |
| | 23Y | 2–3 | 23-yne-1,25-(OH) ₂ D ₃ (10) ^{13a} | |
| | 16E | 2–5 | 16-ene-1,25-(OH) ₂ D ₃ (9) ^{13a,14a} | |
| 18-nor | 18N | 5–10 | 18-nor-1,25-(OH) ₂ D ₃ (11) ¹⁵ | |
| | elongation [homo (H) or di homo (DH)] | 24H | 5–10 | 24-homo-1,25-(OH) ₂ D ₃ (13) ^{16a–c} |
| | | 24DH | 2–5 | 24-dihomo-1,25-(OH) ₂ D ₃ (14) ^{16c} |
| fluorination | 26,27DH | 2–5 | 26,27-dihomo-1,25-(OH) ₂ D ₃ (15) ^{16f,g} | |
| | F2 | 5–10 | 24-F ₂ -1,25-(OH) ₂ D ₃ (16) ^{17a,b} | |
| | F6 | 10 | 26,27-F ₆ -1,25-(OH) ₂ D ₃ (17) ¹⁸ | |
| 20-methyl | 20Me | 7 | 20-Me-1,25-(OH) ₂ D ₃ (38) ¹⁹ | |

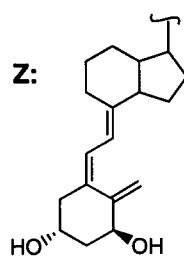
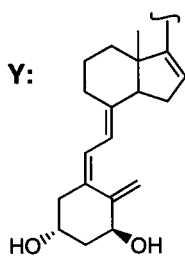
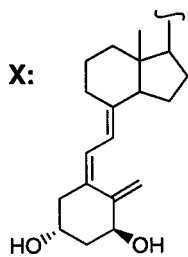
Table 2. Combination of Active Modifications

| entry | modification | compound | cell diff ^a |
|-------|---------------------------|--|------------------------|
| 1 | 24H + 26,27DH | 24,26,27-trihomo-1,25-(OH) ₂ D ₃ (24) ^{8a} | 5 (u) |
| 2 | 16E + 23Y | 16-ene-23-yne-1,25-(OH) ₂ D ₃ (12) ^{13a,14a} | 10 (h) |
| 3 | 22E + 24H | 22-ene-24-homo-1,25-(OH) ₂ D ₃ (20) ^{16a} | 10 (h) |
| 4 | F2 + 24H | 24-F ₂ -24-homo-1,25-(OH) ₂ D ₃ (25) ^{16e,f} | 10 (h) |
| 5 | 22E + 24DH | 22-ene-24,24-dihomo-1,25-(OH) ₂ D ₃ (21) ^{16c} | 10 (h) |
| 6 | 22O + 24H + 26,27DH | 22-oxa-24,26,27-trihomo-1,25-(OH) ₂ D ₃ (19) ^{8a} | 20 (u) |
| 7 | F6 + 22E | 26,27-F ₆ -22-ene-1,25-(OH) ₂ D ₃ (26) ^{13a} | 30 (h) |
| 8 | 22E + 26,27DH | 22-ene-26,27-dihomo-1,25-(OH) ₂ D ₃ (22) ^{13c} | 60 (h) |
| 9 | 22E + 24E + 24H + 26,27DH | 22,24-diene-24,26,27-trihomo-1,25-(OH) ₂ D ₃ (23) ^{13b} | 67 (u) |
| 10 | 16E + 23Y + F6 | 16-ene-23-yne-26,27-F ₆ -1,25-(OH) ₂ D ₃ (27) ^{14b} | 80 (h) |
| 11 | EP + 24H | 20-epi-24-homo-1,25-(OH) ₂ D ₃ (31) ^{8a} | 200 (u) |
| 12 | EP + 24H + 26,27DH | 20-epi-24,26,27-trihomo-1,25-(OH) ₂ D ₃ (32) ^{8a} | 200 (u) |
| 13 | EP + 22E + 24H + 26,27DH | 20-epi-22-ene-24,26,27-trihomo-1,25-(OH) ₂ D ₃ (33) ^{8d} | 1000 (u) |
| 14 | EP + 22O + 24H | 20-epi-22-oxa-24-homo-1,25-(OH) ₂ D ₃ (34) ^{8a} | 1176 (u) |
| 15 | EP + 22O + 24H + 26,27DH | 20-epi-22-oxa-24,26,27-trihomo-1,25-(OH) ₂ D ₃ (35) ^{8a,4d} | 20000 (u) |

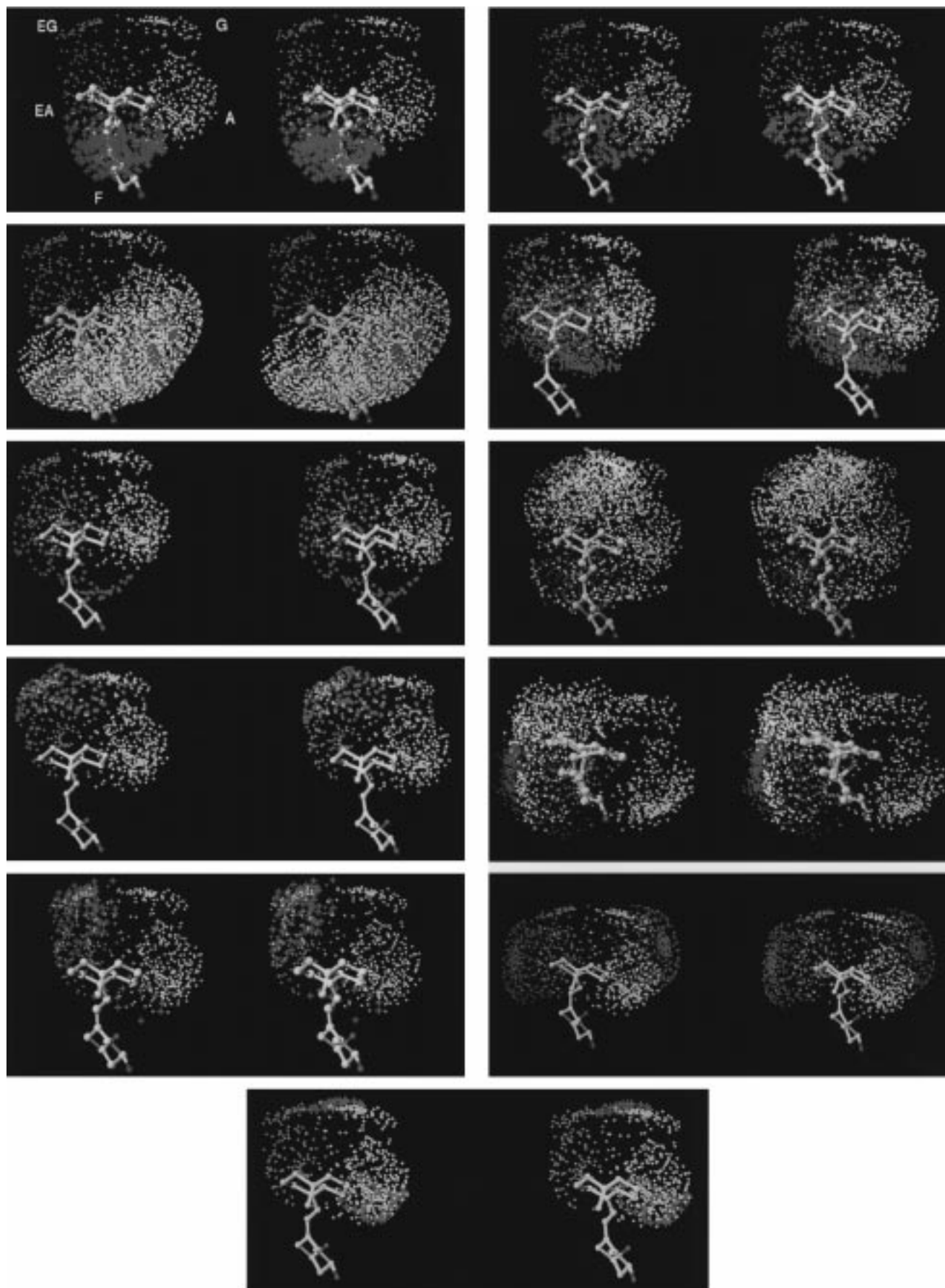
^a Relative differentiating activity of U 937 (u) or HL-60 (h) cells when the reference value of 1,25-(OH)₂D₃ (**1**) is defined as 1.

Chart 1

| | | | | | |
|-----------------|--|-----------------|--|-----------------|--|
| 1 | | 15 | | 29 | |
| 2 | | 16 | | 30 | |
| 3 | | 17 | | 31 ^a | |
| 4 | | 18 ^a | | 32 | |
| 5 | | 19 | | 33 | |
| 6 | | 20 ^a | | 34 ^a | |
| 7 ^a | | 21 | | 35 | |
| 8 ^a | | 22 | | 36 | |
| 9 ^a | | 23 ^a | | 37 ^a | |
| 10 ^a | | 24 | | 38 ^a | |
| 11 ^a | | 25 | | 39 ^a | |
| 12 ^a | | 26 | | 40 ^a | |
| 13 | | 27 | | 41 | |
| 14 | | 28 | | | |



^a The numbered bonds were rotated on the conformational search.



plotted on a dot map (graph sweep in SYBYL) in which the skeleton (the A, seco-B, C, and D ring moieties) is shown with a ball-and-stick model and the terminal oxygen position of each possible conformation is shown as a dot. (4) The mobile areas of the analogue's side chain were compared with the areas A, G, EA, and EG, by superimposing the dot maps of the analogues with those of 1,25-(OH)₂D₃ (**1**) and 20-epi-1,25-(OH)₂D₃ (**2**) at the skeleton.

Results

Compounds with a Side Chain at the F (Front) Region. We have shown that vitamin D compounds are highly potent when the 25-hydroxyl group is located in the A or EA region.^{7a,b} We found a group of compounds in which the 25-oxygen occupies an area in front of the EA region. We call this area F. Compounds which occupy this area have 22-oxa, 16-ene, 22-ene, or 18-nor modifications, such as 22-oxa-1,25-(OH)₂D₃ (**7**),¹² 22-ene-1,25-(OH)₂D₃ (**8**),^{13a} 22,24-diene-24,26,27-trihomo-1,25-(OH)₂D₃ (**23**),^{13b} 16-ene-1,25-(OH)₂D₃ (**9**),^{13a,14a} and 18-nor-1,25-(OH)₂D₃ (**11**).¹⁵

22-Oxa-1,25-(OH)₂D₃ (7). 22-Oxa-1,25-(OH)₂D₃ (**7**) was the first compound in which cell-differentiating and calcemic activities of vitamin D were separated. Compound **7** has about 10-fold higher cell-differentiating activity than 1,25-(OH)₂D₃ (**1**), but it has low calcemic activity.¹² It has not been explained in structural terms why replacing C(22)H₂ with oxygen dramatically enhances potency. Conformational analysis provided one explanation. Replacing C(22)H₂ with oxygen significantly changes the mobility of the 25-oxygen. The 25-oxygen of **7** populates in the F region which lies in front of EA (Figure 2a), whereas the 25-oxygen of 1,25-(OH)₂D₃ (**1**) occupies A and G.

The C(22)H₂ and 22-oxa compounds differ in their minimum-energy conformation. These two compounds differ in their C(16)–C(17)–C(20)–C(22) torsion angle in the minimum-energy conformation. The C(22)H₂ compound adopts a C(16)–C(17)–C(20)–C(22) gauche(+) conformation (Figure 3, yellow), whereas the 22-oxa compound has a gauche(–) conformation (Figure 3, red). In the gauche(–) conformation, C(22)H₂ suffers severe steric repulsion from the C(18)H₃ group. However in the 22-oxa analogue this repulsion is reduced because the oxygen bears no hydrogen, so it adopts a C(16)–C(17)–C(20)–C(22) gauche(–) conformation. When vitamin D adopts a C(16)–C(17)–C(20)–C(22) gauche(+), the 25-oxygen tends to reside in the A and G regions, whereas when it adopts a gauche(–) conformation the 25-oxygen tends to locate in the F region.

22-Dehydro-1,25-(OH)₂D₃ derivatives. Introduction of a π-bond at C(22) similarly reduces the steric repulsion between the C(22) part and the C(18)H₃ group. Therefore, the 25-oxygen of 22-ene-1,25-(OH)₂D₃ (**8**)^{13a} occupies wide areas of the F–A regions (Figure 2b). 22-, 24-Diene-24,26,27-trihomo-1,25-(OH)₂D₃ (**23**),^{13b} which

is undergoing clinical trials as a drug to treat cancer, occupies similar regions (Figure 2c, red crosses). Since this compound has an elongated and rigid conjugated diene structure, the area occupied by its 25-oxygen moves further toward the front and the mobile area is restricted compared with the more flexible 22-ene-24-homo-1,25-(OH)₂D₃ (**20**) (Figure 2c, white dots).

16-Ene-1,25-(OH)₂D₃ Derivatives.¹⁴ The high potency of these vitamin D analogues has been known for a long time, but the reasons were not understood. Introducing a double bond to C(16) also directs the side chain to the front region F. Figure 2d shows that the area occupied by the 25-oxygen of 16-ene-1,25-(OH)₂D₃ (**9**) is similar to that occupied by the 22-oxa (Figure 2a) and 22-ene (Figure 2b) analogues. In the minimum-energy conformation, the calculated C(16)–C(17)–C(20)–C(22) torsion angle of 16-ene-1,25-(OH)₂D₃ (**9**) was –54°. Introducing a triple bond to C(23) and perfluorination at C(26) and C(27) create analogues with increased potency, **12** (Figure 2e) and **27**.

18-Nor-1,25-(OH)₂D₃ (11).¹⁵ Removal of the angular C(18)H₃ group eliminates congestion between the methyl group and C(22)H₂. The side chain therefore can rotate nearly freely around the 17,20 bond. Thus, when all possible conformations were plotted, the area occupied by the 25-oxygen of **11** extended over a wide range (Figure 2f, white dots). However, when 20% stable conformations are selected, the 25-oxygen is highly concentrated in the F region (Figure 2f, red dots). Thus, we explain that this compound has high potency because its side-chain oxygen is located in the active F region.

Compounds with Side Chains in the EA and Left EA (L-EA) Regions. So far the 20-epimerization has the highest effect on increasing vitamin D potency. The HL-60 differentiating activity is enhanced 25–30-fold.⁸ We reasoned that 20-epi-1,25-(OH)₂D₃ (**2**) has high potency because its 25-oxygen can occupy the most active EA region.^{7a} In the minimum-energy conformation, 20-epi-1,25-(OH)₂D₃ analogues adopt the C(16)–C(17)–C(20)–C(22) anti torsion angle (Figure 3, cyan).

20-Epi-16-ene-1,25-(OH)₂D₃ Derivatives. The 20-epimerization of 16-ene analogues has the reverse effect:^{14d,e} 20-Epi-16-ene-23-yne-1,25-(OH)₂D₃ (**37**) is one-half as active as 1,25-(OH)₂D₃ (**1**).^{14d} Therefore, the 20-epimerization of 16-ene-23-yne **12** decreased potency by 1/20. This difference in potency can be understood by comparing the dot map of 16-ene-23-yne-1,25-(OH)₂D₃ (**12**) (Figure 2e) with that of 20-epimer **37** (Figure 2g). While the dots of **12** are found over the active front regions of A, F, and EA, those of **37** are distributed over the inactive rear regions of EG–G.

20-Epi-22-oxa-1,25-(OH)₂D₃ derivatives. We examined the effect of replacing C(22)H₂ with oxygen in the 20-epivitamin D series. Such replacement signifi-

Figure 2. Side-chain mobile area of potent 1,25-(OH)₂D₃ analogues compared with the four regions, A, G, EA, and EG, of **1** (yellow) and **2** (cyan) (stereoviews). Dot maps of each analogue were overlaid with those of **1** and **2**, the skeleton being superimposed: (a, row 1, left) 22-oxa-1,25-(OH)₂D₃ (**7**) (red); (b, row 1, right) 22-ene-1,25-(OH)₂D₃ (**8**) (red); (c, row 2, left) 22,24-diene-24,26,27-trihomo-1,25-(OH)₂D₃ (**23**) (red crosses) and 22-ene-24-homo-1,25-(OH)₂D₃ (**20**) (white dots); (d, row 2, right) 16-ene-1,25-(OH)₂D₃ (**9**) (red); (e, row 3, left) 16-ene-23-yne-1,25-(OH)₂D₃ (**12**) (red); (f, row 3, right) 18-nor-1,25-(OH)₂D₃ (**11**) (white dots, all possible conformations; red dots, 20% stable conformations); (g, row 4, left) 20-epi-16-ene-23-yne-1,25-(OH)₂D₃ (**37**) (red); (h, row 4, right) 20-epi-24-homo-1,25-(OH)₂D₃ (**31**) (white) and 20-epi-22-oxa-24-homo-1,25-(OH)₂D₃ (**34**) (red); (i, row 5, left) 20-methyl-1,25-(OH)₂D₃ (**38**) (red); (j, row 5, right) (17E)- and (17Z)-ene-22-yne-24,26,27-trihomo-1,25-(OH)₂D₃ (**40** and **39**) (right and left red dots, respectively); (k, row 6) 23-yne-1,25-(OH)₂D₃ (**10**) (red).

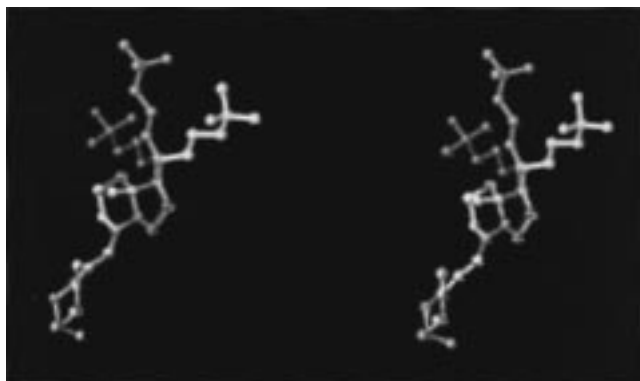


Figure 3. Three staggered conformations at the C(16)–C(17)–C(20)–C(22) torsion angle (stereoview). C(16)–C(17)–C(20)–C(22) gauche(+) (yellow), gauche(–) (red), and anti (cyan) conformations are overlaid.

cantly elevates potency in the 20-epi-24-homovitamin D series. Comparisons of entries 14 and 11 and/or 15 and 12 in Table 2 indicate an activation factor of 5–100 for the 22-oxa modification of 20-epi-24-homovitamins. A difference is also evident in the dot maps: Whereas the dots of the C(22)H₂ compound **31** are distributed over EG and EA (Figure 2h, white dots), EG being more densely populated, those of the 22-oxa analogue **34** are restricted in the left side of EA (L-EA) (Figure 2h, red dots). Their C(17)–C(20)–C(22)–C(23) torsion likely causes these effects. In the minimum-energy conformation, the C(22)H₂ compound **31** adopts a gauche(–) conformation, whereas the 22-oxa analogue **34** adopts an anti torsion angle. The 22-oxygen did not seem to affect the potency of analogues with a side chain of normal length,^{8a} although the trends in the dot maps were similar to those of 24-homo analogues (dot map not shown). The potency of 22-oxa compound **18** was reduced by 1/6.6 relative to that of the C(22)H₂ compound **2**. Since oxygen can form hydrogen bonds and the contribution of a hydrogen bond to a protein–ligand complex can vary depending upon its microscopic environment, the hydrogen-bonding network might alter the effect of the 22-oxygen.

20-Methyl-1,25-(OH)₂D₃. 20-Methylation of 1,25-(OH)₂D₃ elevates the potency:¹⁹ In terms of cell-differentiating activity (HL-60), 20-methyl-1,25-(OH)₂D₃ (**38**) is 7.1 times as active as 1,25-(OH)₂D₃ (**1**). Conformational analysis explains this effect of the 20-methyl group (Figure 2i). This modification makes the three staggered conformations at the C(16)–C(17)–C(20)–C(22) angle of similar energy, the energy increasing in the order anti, gauche(–), and gauche(+). Thus, the dots of **38** are distributed over the EG, EA, and F regions. The dot map suggests that the potency of **38** is between that of **2** and **7**.

17(20)-Dehydrovitamin D. Introduction of a double bond to the 17, 20 position fixes the side-chain direction. In (*Z*)- and (*E*)-17-ene-22-yne-24,26,27-trihomo-1,25-(OH)₂D₃ analogues (**39** and **40**),^{13d} the 25-oxygen atoms are located in opposite directions and can never overlap (Figure 2j). The dots of **39** are located in the L-EA region, while those of **40** are distributed over the right side of the G and A regions. The 17*Z*-isomer **39** is 280-fold more potent than the 17*E*-isomer **40** (potency relative to that of **1** is 710 and 2.5, respectively). Thus, the large potency difference between the two geo-

metrical isomers **39** and **40** can be explained by the difference of the direction of their 25-oxygen.

Other Modifications. 1. Elongation of the Side Chain. Elongation of the side chain increases potency 5–10-fold in both 20-normal^{16a–c} and 20-epivitamin D^{8a} analogues. Side-chain elongation in the 20-normal series expands the area accessible by the 25-oxygen to part of the EA region. This may affect potency. Another and more likely explanation common to both 20-normal and 20-epi compounds is that the 25-oxygen can become closer to the binding site of the ligand-binding domain (LBD) of VDR. If the 25-hydroxyl group of 1,25-(OH)₂D₃ (**1**) binds to LBD through a water molecule, that of the homologues can bind directly and tightly to the same binding site.

2. Entropy Effect by Restriction of Conformation. Introducing a triple bond often elevates potency. This function significantly restricts conformational mobility and thus favorably affects entropy in binding to the receptor. Generally, rigid compounds are more potent than flexible compounds with a similar structure.²¹ The mobility of the side chains of 23-yne-1,25-(OH)₂D₃ (**10**) and 16-ene-23-yne-1,25-(OH)₂D₃ (**12**) is restricted compared with that of 1,25-(OH)₂D₃ (**1**) and 16-ene-1,25-(OH)₂D₃ (**9**), respectively, as shown in Figure 2k,e compared with Figure 2d. In accordance with these figures, the activity of the compounds **10** and **12** is higher than that of the respective compounds **1** and **9**.

Discussion

Since the discovery of active vitamin D₃, several hundred analogues have been synthesized mainly for developing new drugs. Data accumulated from the structure–function studies of the side-chain analogues have revealed nearly 10 modifications that enhance potency (cell differentiation). So far the most activating modification is 20-epimerization. Besides this, 22-oxa, 16-ene, 18-nor, 22-ene, 23-yne, side-chain carbon homologation, perfluorination at the terminal carbons, and 20-methylation also elevate potency. However a systematic theoretical study explaining the relationship between each of these modifications and activity has not been reported.

We have shown, using conformationally restricted analogues (**3–6**), that the area occupied by the 25-hydroxyl group of vitamin D derivatives can be correlated to their affinity for VDR.^{7a} Furthermore, the transcriptional and cell-differentiating activities of these compounds (**3–6**) are demonstrated to be proportional to their VDR affinity.²² We investigated the relationship between the cell-differentiating activity and the conformations of known potent vitamin D analogues based upon our active spatial area concept. We focused upon likely areas occupied by the biologically important 25-hydroxyl group. These areas were searched starting with the minimum-energy conformation, so they were usually found around the minimum-energy conformation of the compound concerned. We occasionally restricted conformations by energy to show more clearly the areas that were more likely to be occupied (as described above).

Our theory successfully correlated the structure and activity of the analogues shown in Tables 1 and 2.²³ We

Table 3. Spatial Region and Activity

| region | VDR affinity ^a | cell diff ^b | representative compound |
|---------|---------------------------|------------------------|---|
| EG | 1/100 (p) 1/250 (b) | ND ^c | 22 <i>S</i> -Me-20-epi-1,25-(OH) ₂ D ₃ (6) ^{7a} |
| G | 1/60 (p) 1/50 (b) | 1/80 (h) | 22 <i>R</i> -Me-1,25-(OH) ₂ D ₃ (3) ^{7a,b} |
| A + G | 1 (p) 1 (b) | 1 (h) | 1,25-(OH) ₂ D ₃ (1) |
| A | 1/3 (p) 1/3 (b) | 1 (h) | 22 <i>S</i> -Me-1,25-(OH) ₂ D ₃ (4) ^{7a,b} |
| F | ND ^c | 10 (h) | 22-oxa-1,25-(OH) ₂ D ₃ (7) ¹² |
| EA + EG | 5 (b) | 25–30 (u) | 20-epi-1,25-(OH) ₂ D ₃ (2) ^{8a} |
| EA | 20 (p) 11 (b) | 200 (h) | 22 <i>R</i> -Me-20-epi-1,25-(OH) ₂ D ₃ (5) ^{7a} |
| L-EA | ND ^c | 1000 (u) | 20-epi-22-oxa-24-homo-1,25-(OH) ₂ D ₃ (34) ^{8a} |

^a Relative VDR affinity determined in our laboratory using porcine intestinal VDR (p) and bovine thymus VDR (b), the reference value of 1,25-(OH)₂D₃ (**1**) being defined as 1. ^b Relative differentiating activity of HL-60 (h) or U 937 (u) cells when the reference value of 1,25-(OH)₂D₃ (**1**) is defined as 1. ^c Not determined in our laboratory.

confirmed that the 25-oxygen of highly potent analogues locates around the EA region. We found that the 25-oxygen of potent 20-normal vitamin D analogues, which have 22-oxa, 16-ene, 22-ene, or 18-nor structures, is found in front of the most active EA region (F) (Figure 2a–f). This finding explains why these compounds are highly potent, although they do not have the 20-epi configuration. We argue that 20-epi-22-oxa-24-homo-vitamin D analogues **34** and **35** have high potency because their side chain is located in the left EA region (Figure 2h). Analogues with a fixed 17,20 bond, **39** and **40**, show that the left regions are more active than the right regions (Figure 2j). Also a comparison of 16-ene-23-yne-1,25-(OH)₂D₃ (**12**) with its 20-epimer (**37**) indicated that the front regions are more potent than the rear regions (Figure 2e,g). The relationship between the activity and 25-oxygen area is summarized in Table 3.

Modifications that do not significantly affect the conformational mobility of the 25-oxygen are side-chain elongation, methylation of the terminal geminal methyl groups, and perfluorination of the terminal methyls and C(24)H₂. Elongation of the side chain increases potency probably because the ligand and the protein can become closer. Conversely, truncating the side chain reduced the potency of 24-nor- (**28**) and 23,24-dinor-1,25-(OH)₂D₃ (**29**) to differentiate HL-60 cells by factors of 1/13 and 1/220, respectively, compared with 1,25-(OH)₂D₃.^{16b} Fluorination and methylation of the terminal positions increase hydrophobicity of the terminal regions and increase the lipophilic contact area, which yields high binding energy.^{21,24} Exposing the 25-hydroxyl group by removing the terminal gem-dimethyl groups lowers the cell-differentiating activity of 26,27-dinor-1,25-(OH)₂D₃ (**30**) 100-fold relative to 1,25-(OH)₂D₃.^{16b}

The results of the present study showed that the biological activity of vitamin D derivatives is correlated with the area occupied by the 25-oxygen. Since this area is found around the minimum-energy conformation of vitamin D, these results indicate that the potency of vitamin D derivatives is correlated with their minimum-energy conformations. It is generally accepted that a ligand binds more tightly to the receptor when the ligand adopts a more stable conformation.^{11,21} We confirmed that this is also true of vitamin D analogues.

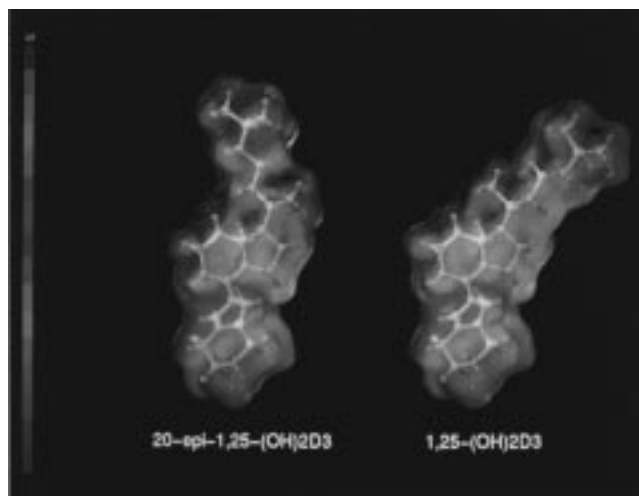


Figure 4. Surface structures of 1,25-(OH)₂D₃ (**1**) and 20-epi-1,25-(OH)₂D₃ (**2**) in which the side chains are directed to the A and EA regions, respectively. Colors show lipophilic properties, brown and blue being the colors of the most and the least lipophilic nature, respectively.

These studies also allowed us to image a topological feature of the LBD of VDR facing the 25-hydroxyl group of the ligand. Figure 4 shows the surface structures of 1,25-(OH)₂D₃ (**1**) and 20-epi-1,25-(OH)₂D₃ (**2**) in which the side chains are located in the A and EA regions, respectively (colors show lipophilic properties, brown and blue being the colors of the most and the least lipophilic nature, respectively).

In conclusion, we showed that the conformational search of vitamin D side chains helps predict the potency of vitamin D analogues. A similar approach would be useful for predicting the potency of analogues with seco-B and A ring modifications. These studies may also be used to discriminate the various activities of vitamin D, although this would require further study.

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