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Chemical synthesis, biological activities and action on nuclear receptors of 20*S*(OH)D₃, 20*S*,25(OH)₂D₃, 20*S*,23*S*(OH)₂D₃ and 20*S*,23*R*(OH)₂D₃

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Abstract

New and more efficient routes of chemical synthesis of vitamin D₃ (D₃) hydroxy (OH) metabolites, including 20*S*(OH)D₃, 20*S*,23*S*(OH)₂D₃ and 20*S*,25(OH)₂D₃, that are endogenously produced in the human body by CYP11A1, and of 20*S*,23*R*(OH)₂D₃ were established. The biological evaluation showed that these compounds exhibited similar properties to each other regarding inhibition of cell proliferation and induction of cell differentiation but with subtle and quantitative differences. They showed both overlapping and differential effects on T-cell immune activity. They also showed similar interactions with nuclear receptors with all secosteroids activating vitamin D and aryl hydrocarbon receptors in functional assays and also as indicated by molecular modeling. They functioned as substrates for CYP27B1 with enzymatic activity being the highest towards 20*S*,25(OH)₂D₃ and the lowest towards 20*S*(OH)D₃. In conclusion, defining new routes for large scale synthesis of endogenously produced D₃-hydroxy derivatives by

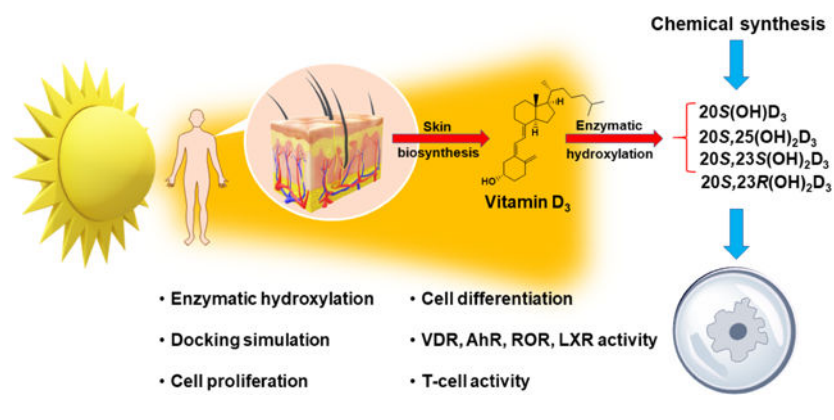
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pathways initiated by CYP11A1 opens an exciting era to analyze their common and differential activities *in vivo*, particularly on the immune system and inflammatory diseases.

Graphical Abstract



Introduction

Vitamin D₃ (**1**, Fig. 1) is formed from 7-dehydrocholesterol (7DHC) by the action of ultraviolet (UVB) radiation [1–4]. The electrocyclic B-ring opening generates previtamin D₃ which is converted to vitamin D₃ by a first-order thermal isomerization in the skin [1–3]. Vitamin D₃ is activated by hydroxylation at C-25 by CYP2R1 or CYP27A1 then at C1 α by CYP27B1, to produce 1 α ,25(OH)₂D₃ (**2**) [4–7]. In addition to regulating calcium homeostasis, 1 α ,25(OH)₂D₃ has pleiotropic effects affecting almost all body functions through an interaction with vitamin D receptor (VDR) [4, 5, 8–16]. 1 α ,25(OH)₂D₃ regulates cell proliferation and differentiation, exhibits anticancer properties, regulates skin functions, inhibits adaptive immunity, and upregulates innate immune responses among other physiological functions [10, 17–21]. It is clinically used to treat autoimmune and inflammatory diseases, including various skin disorders [22–31]. However, toxic/calcemic effects of 1 α ,25(OH)₂D₃ at pharmacological doses constrains its use to treat autoimmune diseases and other pathological states [10, 12, 32–34].

New pathways of vitamin D₃ activation by CYP11A1 producing many hydroxy-derivatives have now been well characterized [7, 35]. The main sequence of the reactions starts with hydroxylation at C-20 to produce 20S(OH)D₃ (**3**) which is sequentially hydroxylated to 20S,23S(OH)₂D₃ (**4a**) and 17R,20S,23S(OH)₃D₃ (**5**) [35–37]. 20S(OH)D₃ can also be hydroxylated by CYP27A1, CYP24A1 and CYP2R1 to produce 20S,25(OH)₂D₃ (**6**), which is an excellent substrate for CYP27B1 [7]. These metabolites are produced *ex vivo* in skin cells, placenta and adrenal glands in a CYP11A1 dependent fashion [38, 39]. Some, including 20S(OH)D₃ and 1 α ,20S(OH)₂D₃, have been measured in the nM range in all human serum samples tested [40, 41] and 20S(OH)D₃ was recently detected in honey [42]. Thus, in addition to functioning as hormones in humans (see below), some of these secosteroids can be defined as natural products [43].

In vitro studies have demonstrated that 20 S (OH)D₃ and 20 S ,23 S (OH)₂D₃ show anti-proliferative, pro-differentiation, anticancer, photoprotective, anti-inflammatory and anti-fibrinogenic properties [44–52]. 20 S ,25(OH)₂D₃ has also been reported to exhibit anti-melanoma activity in cell culture [53]. Anti-fibrogenic, photoprotective and anti-melanoma properties for 20 S (OH)D₃ were also demonstrated in *in vivo* models [50, 54, 55]. Of high clinical relevance is that CYP11A-derived 20 S (OH)D₃ and 20 S ,23 S (OH)₂D₃ are non-calcemic at doses that are highly toxic for 1 α ,25(OH)₂D₃ and 25(OH)D₃ [32, 50, 56]. The CYP11A1-derived secosteroids have been shown to mediate their effects by acting as biased agonists on the VDR [51, 57–59], inverse agonists on retinoic acid orphan receptors (ROR) α and γ [58, 60], and agonists on liver X receptors (LXRs) [61]. 20 S (OH)D₃ and 20 S ,23 S (OH)₂D₃ are also agonists for the aryl hydrocarbon receptor (AhR) [62].

To provide a robust source of key CYP11A1-derived secosteroids produced by pathways initiated by CYP11A1, and to enable better characterization of their biological activity, we describe new chemical routes for large scale synthesis of 20 S (OH)D₃ (**3**), 20 S ,23 S (OH)₂D₃ (**4a**), 20 S ,23 R (OH)₂D₃ (**4b**) and 20 S ,25(OH)₂D₃ (**6**).

Results and Discussion

Chemical synthesis and confirmation of the structures

Continuing our studies related to vitamin D₃ derivatives we designed an efficient synthetic route for 20 S (OH)D₃ and its 23- as well as 25-hydroxylated analogs 20 S ,23 S (OH)₂D₃, 20 S ,23 R (OH)₂D₃, 20 S ,25(OH)₂D₃. Convergent synthesis was used for the preparation of these target compounds where the key stage was a Wittig-Horner coupling of the corresponding Grundmann ketone (CD-ring fragment) with A-ring phosphine oxide. Starting from vitamin D₂ (**7**), the required building blocks **8** – **12** (Fig. 2) were prepared using a slight modification of literature procedures; the synthetic steps (Scheme S1), procedures and characterization of the compounds obtained are presented in Supporting Information.

The ketone **12** was treated with two different Grignard reagents [63, 64] providing diastereomerically pure 20 S -diols **13** and **14** (Scheme 1). Secondary hydroxyl groups in **13** and **14** were efficiently oxidized to the corresponding ketones **15** and **16** with tetrapropylammonium perruthenate in the presence of *N*-methylmorpholine-*N*-oxide. Tertiary hydroxyls were then protected as TMS ethers providing CD-ring building blocks **17** and **18**. These hydrindanones were then coupled with the known A-ring phosphine oxide **9** [65] to furnish protected vitamin D₃ derivatives **19** and **20**. Their deprotection followed by RP-HPLC purification provided the target 20-hydroxy compounds **3** and **6** in good yields and high purity.

Methyl ketone **11** was then used for the preparation of desired 20 S ,23 S (OH)₂D₃ as well as its epimer 20 S ,23 R (OH)₂D₃ (Scheme 2). As a consequence of the steric effect of both 7a'-methyl group and 4'-OTES substituents, only the 3 S -epimer **21** was isolated when the ketone **11** was treated with the nitrile anion, generated from acetonitrile and LDA. The analogous, highly stereoselective addition of Grignard reagents has been thoroughly studied by us previously [63,64]. The secondary hydroxyl group in **21** was deprotected and the

resulting hydrindanol **22** was subjected to the oxidation with TPAP/NMO leading to the ketone **23**. Protection of the tertiary hydroxyl group in this compound by its treatment with trimethylsilyl triflate provided the desired building block **24**. Wittig-Horner olefination was used as a coupling reaction of the hydrindanone **24** and phosphine oxide **9**. The formed nitrile **25**, possessing the vitamin D skeleton, was reduced with diisobutylaluminum hydride to the corresponding aldehyde **26**. In this compound elongation of the side chain was effected by its Grignard reaction with *i*-butyl magnesium chloride providing a mixture of diastereomeric alcohols **27a,b**. Finally, removal of silyl protecting groups and efficient separation by high-performance liquid chromatography (HPLC) provided the target epimeric (20*S*,23*S*)- and (20*S*,23*R*)-dihydroxylated vitamin D₃ compounds (**4a,b**).

The above scheme represents a significant improvement to the synthetic route for the preparation of 20*S*(OH)D₃ compared to the method described by Li et al [65]. The advantage of the synthesis reported here is the application of a different protecting group of the tertiary hydroxyl at C-20. Thus, the use of trimethylsilyl ether (OTMS), being significantly more labile than EOM ether, allowed us to obtain the final vitamin D compound with significantly increased (ca. fourfold) overall yield. Moreover, using analogous conditions, 20*S*,25(OH)₂D₃ has also been successfully prepared, representing another active metabolite of vitamin D₃, that so far has only been obtained enzymatically [66].

The synthesis of 23-hydroxylated derivatives presented above has also been significantly improved in comparison with the previously reported synthetic path [67] involving a series of linear transformations. The epimeric 20,23-dihydroxyvitamin D₃ metabolites were obtained in high yield, easily separated by HPLC and individual compounds were biologically evaluated. The synthetic strategy applied by us gave a marked (ca. 5 times) increase in the overall yield of the target secosteroids, avoiding the irradiation step used in past routes which limits the scale of the conversion of the 7-dehydrocholesterol analog to the corresponding secosteroid. Moreover, almost all steps are executed on the optically pure compounds; formation of C-23 stereoisomers takes place at the final stage of our synthesis.

Hydroxylation at C1 α

All four chemically synthesized secosteroids were converted to their expected 1 α -hydroxy derivatives by CYP27B1, and identified on the basis of previous studies on the enzymatically and/or chemically derived substrates [63, 66–68]. The rates varied dramatically with a high rate of 1 α -hydroxylation being observed for 20*S*,25(OH)₂D₃ (Table 1), consistent with previous reports for this substrate produced enzymatically [66]. The naturally occurring diastereomer of 20,23(OH)₂D₃, the 20*S*,23*S*-isomer, was metabolized approximately twice as fast as the non-natural 20*S*,23*R*-isomer (Table 1), as noted in a previous study [67]. 20*S*(OH)D₃ was a relatively poorer substrate for CYP27B1 compared to 20*S*,25(OH)₂D₃ (Table 1), also consistent with a previous report [66]. In summary, the efficiency of metabolism of D₃ derivatives by CYP27B1 is as follows: 20*S*,25(OH)₂D₃ >> 20*S*,23*S*(OH)₂D₃ > 20*S*(OH)D₃ = 20*S*,23*R*(OH)₂D₃.

Regulation of cell proliferation and differentiation program

The anti-proliferative properties of $20S(OH)D_3$ and $20S,23(OH)_2D_3$ [44, 48–50, 52, 55, 56, 69–72] and to some degree of $20S,25(OH)_2D_3$ [53, 58] are well established. We tested the capabilities of the newly synthesized secosteroids to inhibit proliferation and found that $20S,25(OH)_2D_3$ inhibits fibroblast and keratinocyte proliferation in similar manner to its precursor, $20S(OH)D_3$ (Fig. 3). Similar effects were also observed for both $20S,23(OH)_2D_3$ isomers (Fig. 3), which is consistent with previously reported data [67]. Again, analysis of human keratinocytes showed that $20S,25(OH)_2D_3$ was slightly better at inducing gene expression of involucrin, CK10, and filaggrin, indicative of keratinocyte differentiation, compared to $20S(OH)D_3$ (Fig. 4A) with $20S,23R(OH)_2D_3$ and $20S,25(OH)_2D_3$ showing similar effects on the expression of these genes (Fig. 4B). At the protein level, all compounds including $20S(OH)D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$ and $20S,25(OH)_2D_3$ showed similar net biological effects in increasing the levels of involucrin, CK10, CK14, and catalase (Fig. 4C). These results are consistent with previously published pro-differentiation properties of $20S(OH)D_3$ and $20S,23S(OH)_2D_3$ [32, 45, 48, 49, 69, 70, 73], which we now extend to include $20S,25(OH)_2D_3$ and $20S,23R(OH)_2D_3$.

Nuclear receptor signaling

To better define the properties of the newly synthesized secosteroids, we tested their activities on the VDR, AhR and LXR by performing functional assays which were further validated by qPCR of downstream enzymes (Fig. 5) and molecular modeling (Table 2 and Fig. 6). Binding affinities of top-scored docking complexes of vitamin D₃ hydroxy derivatives with VDR, AhR, LXRs (LXR α , LXR β) and RORs (ROR α , ROR γ) from molecular docking are presented in Table 2, known ligands for each receptor were used as controls. $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ have favorable energies for binding to the ligand binding domain of VDR, AhR, LXRs (LXR α , LXR β) and RORs (ROR α , ROR γ), and are comparable to the binding energies of known ligands for each receptor, including $1\alpha,25(OH)_2D_3$.

All of the newly-synthesised compounds induced VDR translocation to the nucleus (Fig. 5A) in a manner similar to that reported previously for some of these secosteroids [57, 74]. *CYP24A1* is a downstream target of the activated VDR and we observed that $20S,25(OH)_2D_3$ caused much greater stimulation of *CYP24A1* expression than $20S(OH)D_3$. There was no significant difference in the expression of *CYP24A1* caused by $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ but both of these caused significantly less expression than $1\alpha,25(OH)_2D_3$ (Fig. 5C). The results of these assays are supported by molecular docking results of these secosteroids with the VDR (Table 2 and Supplemental S72). $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ all have favorable to binding energies for interaction with VDR (Table 2) and bind to the same LBD of VDR (Supplemental S72).

In summary, these data and analyses are consistent with our previous studies on the VDR which showed that $20S(OH)D_3$ and $20S,23(OH)_2D_3$ act on the genomic site of the VDR with lower efficiency than their 1α -hydroxy derivatives, as a consequence of their different interactions with amino acids of the LBD [48, 57, 59, 67, 74]. The better efficiency

of $20S,25(OH)_2D_3$ in stimulating the expression of *CYP24A1* in comparison to other secosteroids could be secondary to its very efficient hydroxylation by CYP27B1 (Table 1) [66].

We also found that the newly-synthesized secosteroids interacted with the AhR (Fig. 5B, D, Table 2), similar to our previous study [62] except that $20S,23R(OH)_2D_3$ was found to act as a reverse agonist. Molecular modeling predicts that $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ share the same binding region in AhR. $20S(OH)D_3$ and $20S,25(OH)_2D_3$ have the similar binding postures with AhR, while the binding postures of $20S,23R(OH)_2D_3$ and $20S,23S(OH)_2D_3$ with AhR are very similar (Fig. 6a).

The binding between AhR and D_3 derivatives is mainly through hydrophobic interactions. Hydrophobic residues Cys333, Phe295, Phe287, Tyr322, Leu308, Tyr310, Leu315, Leu353, Ile325, Val381 and Ala367 in AhR are directly involved in the binding with all four molecules of $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ (Fig. 6a (B)). Recently we have reported that $20S(OH)D_3$, $20S,23(OH)_2D_3$, $20S,25(OH)_2D_3$ and other D_3 -hydroxy derivatives can act as ligands on LXR [61]. Therefore, we compared interactions of $20S,23S(OH)_2D_3$ versus $20S,23R(OH)_2D_3$ with the LBD of LXR α and β (Fig. 5E, F) using a functional assay and molecular modeling (Table 2, Fig. 6b). The functional assays using LanthaScreen TR-FRET LXR α and β coactivator assays has confirmed that $20S,23(OH)_2D_3$ isomers are agonists on the LXR α and β [61], with some differences (Fig. 5E, F). Vitamin D_3 derivatives had better affinity on LXR β than on LXR α . When binding with LXRs, $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ all bind to the same binding pocket of either LXR α or LXR β (Fig. 6b (A) and (B)). For LXR α , $20S(OH)D_3$ and $20S,25(OH)_2D_3$ have the similar binding postures, whereas $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ have the similar binding posture but twisted from C20 compared to that of $20S(OH)D_3$ and $20S,25(OH)_2D_3$ (Fig. 6b (A) and (C)). For LXR β , similar observation based on 2D and 3D interaction maps (Fig. 6b (B) and (D)). $20S,23S(OH)_2D_3$ has a better affinity with LXR β compared to that of $20S,23R(OH)_2D_3$ (Table 2), which is also observed in experimental data (Fig. 5E, F). $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ binding postures to LXR α (Fig. 6b (C)) are very similar to their binding postures to LXR β (Fig. 6b (D)).

The binding between LXR α and D_3 derivatives is mainly through hydrophobic interactions. Hydrophobic residues Ile295, Phe335, Met298, Leu299, Phe315, Leu316, Ala261, Leu260, Phe257 and Leu439 in LXR α are directly involved in the binding with all four molecules of $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ (Fig. 6b (C)). The binding of $20S(OH)D_3$ to LXR α involved the formation of a hydrogen bond between the hydroxyl group on C3 of $20S(OH)D_3$ with Leu316, while the binding of $20S,23S(OH)_2D_3$ with LXR β involved formation of a hydrogen bond between the 23-hydroxyl group of $20S,23S(OH)_2D_3$ and Leu274 (Fig. 6b (D)).

Lastly, we performed molecular modeling of $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ with LBD of the RORs (Fig. 6c, Table 2). $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$ and $20S,23R(OH)_2D_3$ all bind to the same binding pocket of both ROR α and ROR γ (Fig. 6c (A) and (B)). For ROR α ,

20*S*(OH)₂D₃ and 20*S*,25(OH)₂D₃ share a similar binding posture to each other, distinct from 20*S*,23*S*(OH)₂D₃ and 20*S*,23*R*(OH)₂D₃ which are also similar to each other (Fig. 6c (C)). For ROR γ , again 20*S*(OH)₂D₃ and 20*S*,25(OH)₂D₃ show similar binding postures to each other, as do 20*S*,23*R*(OH)₂D₃ and 20*S*,23*S*(OH)₂D₃ (Fig. 6c (D)). The bindings between RORs and D₃ derivatives are mainly mediated through hydrophobic interactions. Hydrophobic residues Ile327, Tyr290, Met368, Val379, Tyr380, Val364, Val403, Phe391 and Ile400 in ROR α are directly involved in the binding with all four D₃ derivatives (Fig. 6c (C)). Hydrophobic residues Leu287, Phe378, Phe377, Val376, Ile397, Ile400, Phe388, Leu391, Cys320 and Leu324 in ROR γ are directly involved in the binding with all four molecules of 20*S*(OH)₂D₃, 20*S*,25(OH)₂D₃, 20*S*,23*S*(OH)₂D₃ and 20*S*,23*R*(OH)₂D₃ (Fig. 6c (D)). These modeling data not only support our previous studies identifying novel hydroxy derivatives of D₃ as inverse agonists on RORs [58, 60], but also extend such observations to include 20*S*,23*R*(OH)₂D₃ and 20*S*,25(OH)₂D₃.

In summary, chemically synthesized 20*S*(OH)₂D₃, 20*S*,23*S*(OH)₂D₃, 20*S*,23*R*(OH)₂D₃ and 20*S*,25(OH)₂D₃ showed the ability to act on the VDR, AhR, LXR and ROR, consistent with our previous reports [57–62, 75], with some differences as indicated above. These studies provide a background for future exciting studies on the exact nature of these interactions between the secosteroids and their receptors, and the differences between different analogs.

Different vitamin D compounds have distinct and overlapping effects on activation of T cells in human peripheral blood

To determine if different vitamin D₃ compounds variably alter activation human CD4 and CD8 T cell, PBMC from healthy individuals were stimulated with anti-CD3 and anti-CD28 to activate T cells in the absence (ethanol) or presence of 20*S*(OH)₂D₃, 20*S*,23*S*(OH)₂D₃, 20*S*,23*R*(OH)₂D₃, 20*S*,25(OH)₂D₃ or 1 α ,25(OH)₂D₃ for 24 h. The cells were stained for the expression of lineage markers (CD4, CD8, CD14, CD19) and activation induced markers (AIM; CD69, ICOS, OX40, CD137, PD1 and HLADR) [76, 77]. We analyzed the data using an unsupervised algorithm-based approach combining TriMAP followed by self-organizing map (FlowSOM) cluster overlay as described in the Experimental section (Fig. 7A and 7B). Unsupervised algorithm-based analyses of multi-dimensional data enables unbiased identification of discovery as well as changes in cell populations [78,79]. Based on the expression levels of CD4, CD8 and the AIM, we identified four populations each of CD4 and CD8 T cells. Populations 2 and 4 within CD4 T cells were differentially altered by the vitamin D₃ compounds (Fig. 7A). For example, compared to ethanol treated cells, 1 α ,25(OH)₂D₃, 20*S*,23*S*(OH)₂D₃ and 20*S*,25(OH)₂D₃ treated cells contained fewer cells that down modulated CD4 and expressed increased levels of CD69 and ICOS (population 4, Fig. 7B). In contrast, 20*S*(OH)₂D₃ treatment increased the proportion of cells with population 4 phenotype and 20*S*,23(OH)₂D₃ had no effect on this population. Within CD8 T cells population 6 (cells that express ICOS) were differentially modulated by the vitamin D₃ compounds compared to ethanol treated cells (Fig 7A and 7B). . The unbiased algorithm-based computational analysis which used statistical “R code” reveals with accuracy that the different vitamin D compounds have overlapping and distinct effects in their ability to modulate T cell activation.

Concluding remarks

Efficient chemical synthetic routes for $20S(OH)D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$, and $20S,25(OH)_2D_3$ were established. Convergent synthesis was used for the preparation of these target compounds where the key stage was a Wittig-Horner coupling of the corresponding Grundmann ketone (CD-ring fragment) with A-ring phosphine oxide. The resulting secosteroids were substrates for 1α -hydroxylation by CYP27B1 with the following selectivity: $20S,25(OH)_2D_3 \gg 20S,23S(OH)_2D_3 > 20S(OH)D_3 = 20S,23R(OH)_2D_3$. Functional assays and molecular modeling have shown that they act on the VDR, AhR, LXR and ROR α and γ with some notable differences. $20S,25(OH)_2D_3$ showed stronger stimulation of *CYP24A1*, a gene downstream of the VDR, than the other secosteroids which may be explained by its more efficient hydroxylation by CYP27B1. On the other hand, there was an interesting change from agonistic activity of $20S,23S(OH)_2D_3$ on the AhR to inverse agonism by its 23*R*-isomer ($20S,23R(OH)_2D_3$). In addition, $20S,23R(OH)_2D_3$ was less efficient in activation of LXR β than $20S,23S(OH)_2D_3$. This work provides the background for future studies on the different modes of nuclear receptor signaling by hydroxy- D_3 derivatives under study. Interestingly, the major phenotypic effects caused by the synthesized secosteroids were very similar and included inhibition of cell proliferation in skin cells and induction of keratinocyte differentiation. Finally, these secosteroids showed immunomodulatory effects on human PBMC stimulated by coactivating peptides with net effects of downregulation of proinflammatory responses. Interestingly, the pattern of expression of lymphocytic markers was both overlapping and differential indicating a difference in mechanism of action depending on the position of the OH group and its configuration in the D_3 side chain. These secosteroids showed differential effects in comparison to classical $1\alpha,25(OH)_2D_3$. This work underpins new possibilities for exciting studies on the anti-inflammatory actions of the D_3 hydroxy derivatives in order to establish efficient methods to treat inflammatory and autoimmune diseases.

In summary, new efficient routes of chemical synthesis for $20S(OH)D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$, and $20S,25(OH)_2D_3$ were designed allowing their production for in vivo testing and further clarification of different mechanisms of action as well as future preclinical studies on autoimmune disorders including lupus erythematosus, rheumatoid arthritis, inflammatory bowel disease, psoriasis and multiple sclerosis.

Experimental Section

Chemistry

Optical rotations were measured in chloroform using a Perkin-Elmer model 343 polarimeter (Shelton, CT, USA). Ultraviolet (UV) absorption spectra were obtained on a Shimadzu UV-1800 spectrophotometer (Kyoto, Japan) in absolute ethanol. Nuclear magnetic resonance spectra were recorded in $CDCl_3$ solutions using Bruker AVANCE 300 MHz (Karlsruhe, Germany) and Bruker AVANCE 500 MHz instruments. Chemical shifts (δ) are reported in parts per million relative to $(CH_3)_4Si$ (δ 0.00) or the solvent signal as an internal standard. Abbreviations used are singlet (s), doublet (d), triplet (t), quartet (q), multiplet (m), broad (br), narrow (narr). High-resolution mass spectra were recorded on Shimadzu LCMS-IT-TOF mass spectrometer using electrospray ionization (ESI) technique.

Reactions were carried out with magnetic stirring. All reactions involving moisture- or oxygen-sensitive compounds were carried out under dry argon atmosphere. Reaction temperatures refer to external bath temperatures. Tetrahydrofuran was distilled from Na/benzophenone; dichloromethane was distilled from P₂O₅. The organic extracts were dried over anhydrous MgSO₄, filtered and concentrated using a rotary evaporator with vacuum pump attached. Reactions were monitored by thin-layer chromatography (TLC) using aluminum-backed MERCK 60 silica gel plates (Darmstadt, Germany) (0.2 mm thickness). The chromatograms were visualized first with ultraviolet light (254 nm) and then by immersion in a cerium-molybdenum solution [10 g Ce(SO₄)₂ × 4 H₂O, 25 g phosphomolybdic acid, 60 mL H₂SO₄ and 940 mL H₂O] or *p*-anisaldehyde solution (5 mL H₂SO₄, 1.5 mL glacial AcOH, 3.7 mL *p*-anisaldehyde, 135 mL H₂O) followed by heating.

The purity of final compounds was determined by HPLC and they were judged at least 98% pure. For this purpose, straight-phase (SP) and reversed-phase (RP) high-performance liquid chromatography were performed on Shimadzu Prominence system equipped with SPD 20A UV detector (265 nm). For straight-phase a Zorbax RX-SIL semi-preparative 9.4 × 250 mm, 5 μm column and for reverse-phase a Zorbax Eclipse XDB-C18 semi-preparative 9.4 × 250 mm, 5 μm HPLC column were used. The purity and identity of the synthesized secosteroids were additionally confirmed by inspection of their ¹H NMR, ¹³C NMR and high-resolution mass spectra (HRMS).

(1S,3aR,4S,7aS)-1-[(S)-2'-Hydroxy-6'-methylheptan-2'-yl]-7a-methyloctahydro-1H-inden-4-ol (13).—To a suspension

of activated magnesium turnings (105 mg, 4.40 mmol) in anhydrous THF (7 mL) was added a small crystal of iodine and 1-bromo-4-methylpentane (0.22 mL, 1.48 mmol). The mixture was heated to the boiling point and second portion of 1-bromo-4-methyl-pentane (0.22 mL, 1.48 mmol) was added. Stirring was continued for 1 h and then the mixture was cooled to 0 °C. The generated Grignard reagent was then added to a solution of ketone **12** (70 mg, 0.295 mmol) in anhydrous THF (3 mL) at 0 °C and the stirring was continued for 2 h at this temperature and overnight at room temperature. The mixture was diluted with diethyl ether (10 mL) and saturated solution of NH₄Cl (10 mL) was carefully added at 0 °C. The layers were separated and the aqueous was extracted with AcOEt (2 × 10 mL). The combined organic solvents were dried (MgSO₄) and concentrated. The residue was purified by column chromatography on silica using hexane/ethyl acetate (84:16) to give the compound **13** (70 mg, 85%) as a colorless oil. **13**: [α]_D²⁴+21 (c 0.6, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.88 (6H, d, *J* = 6.6 Hz, 6'-CH₃ and 7'-H₃), 1.15 (3H, s, 7a-CH₃), 1.27 (3H, s, 1'-H₃), 4.10 (1H, narr m, 4α-H); ¹³C NMR (75 MHz) δ 15.4, 17.4, 21.3, 21.9, 22.2, 22.5, 22.7, 26.3, 27.9, 33.4, 39.6, 40.8, 42.5, 44.1, 52.6, 57.9, 69.4, 75.1; HRMS (ESI) calcd. for C₁₈H₃₄O₂Na (M⁺+ Na) 305.2457, measured 305.2462.

(1S,3aR,4S,7aS)-1-[(S)-2'-Hydroxy-6'-methyl-6'-[(triethylsilyl)oxy]heptan-2'-yl]-7a-methyloctahydro-1H-inden-4-ol (14).—The reaction of the ketone **12** (90 mg, 0.377 mmol) with the Grignard reagent, generated from 1-bromo-4-methyl-4-[(triethylsilyl)oxy]pentane, was performed as described above for **13**. After the analogous procedure, the crude product was purified by column chromatography on silica

using hexane/ethyl acetate (86:14) to give **14** (124 mg, 85%) as a colorless oil. **14**: $[\alpha]_{\text{D}}^{24} +16$ (*c* 0.95, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.56 (6H, q, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 0.94 (9H, t, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 1.14 (3H, s, 7a-CH₃), 1.19 (6H, s, 6'-CH₃ and 7'-H₃), 1.27 (3H, s, 1'-H₃), 4.09 (1H, narr m, 4a-H); ¹³C NMR (75 MHz) δ 6.8, 7.1, 15.4, 17.4, 19.0, 21.4, 22.2, 26.3, 29.8, 30.1, 33.4, 40.8, 42.6, 44.4, 45.6, 52.6, 57.9, 69.5, 73.4, 75.2; HRMS (ESI) calcd. for C₂₄H₄₈O₃SiNa (M⁺ + Na) 435.3270, measured 435.3279.

(1S,3aR,7aR)-1-[(S)-2'-Hydroxy-6'-methylheptan-2'-yl]-7a-methyloctahydro-4H-inden-4-one (15).—Molecular sieves 4 Å (70 mg) were added to a solution of the diol **13** (70 mg, 0.247 mmol) in anhydrous CH₂Cl₂ (2 mL) and the mixture was cooled to 0 °C. 4-Methylmorpholine *N*-oxide (NMO, 44 mg, 0.371 mmol) and tetrapropylammonium perruthenate (TPAP, 44 mg, 0.124 mmol) were then added. The resulted dark mixture was stirred for 1.5 h and then filtered through Celite. The solvent was evaporated and the residue was purified by column chromatography on silica using hexane/ethyl acetate (86:14) to give the ketone **15** (53 mg, 77%) as a colorless oil. **15**: $[\alpha]_{\text{D}}^{24} -21$ (*c* 1.5, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.83 (3H, s, 7a-CH₃), 0.89 (6H, d, *J* = 6.6 Hz, 6'-CH₃ and 7'-H₃), 1.31 (3H, s, 1'-H₃), 2.45 (1H, dd, *J* = 10.7, 7.0 Hz, 3a-H); ¹³C NMR (75 MHz) δ 14.4, 18.8, 21.8, 22.0, 22.6, 22.7, 23.9, 25.6, 27.9, 39.3, 39.5, 40.8, 44.1, 49.8, 58.2, 62.2, 74.6, 211.9; HRMS (ESI) calcd. for C₁₈H₃₂O₂Na (M⁺ + Na) 303.2300, measured 303.2295.

(1S,3aR,7aR)-1-[(S)-2'-Hydroxy-6'-methyl-6'-[(triethylsilyl)oxy]heptan-2'-yl]-7a-methyloctahydro-4H-inden-4-one (16).—The oxidation of the secondary hydroxyl group in **14** (145 mg, 0.353 mmol) was performed analogously as described above for **13**. The crude product was purified by column chromatography over silica using hexane/ethyl acetate (88:12) to give the ketone **16** (115 mg, 80%) as a colorless oil. **16**: $[\alpha]_{\text{D}}^{24} -13$ (*c* 1.0, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.56 (6H, q, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 0.81 (3H, s, 7a-CH₃), 0.94 (9H, t, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 1.20 (6H, s, 6'-CH₃ and 7'-H₃), 1.30 (3H, s, 1'-H₃), 2.42 (1H, dd, *J* = 10.5, 6.9 Hz, 3a-H); ¹³C NMR (75 MHz) δ 6.8, 7.1, 14.4, 18.8, 19.0, 21.8, 24.0, 26.6, 29.8, 30.1, 39.3, 40.8, 44.4, 45.5, 49.8, 58.1, 62.2, 73.3, 74.7, 211.9; HRMS (ESI) calcd. for C₂₄H₄₆O₃SiNa (M⁺ + Na) 433.3114, measured 433.3122.

(1S,3aR,7aR)-7a-Methyl-1-[(S)-6'-methyl-2'-[(trimethylsilyl)oxy]heptan-2'-yl]octahydro-4H-inden-4-one (17).—To a solution of hydroxy ketone **15** (76 mg, 0.271 mmol) in anhydrous CH₂Cl₂ (2 mL) was added 2,6-lutidine (63 μL, 0.542 mmol) and trimethylsilyl trifluoromethanesulfonate (74 μL, 0.406 mmol) at -78 °C and the mixture was stirred at this temperature for 1 h. Saturated NaHCO₃ (2 mL) was added and the mixture was extracted with dichloromethane (3 × 5 mL). The combined organic layers were washed with 5% HCl (5 mL), water (5 mL), dried (MgSO₄) and evaporated. The residue was purified by column chromatography on silica using hexane/ethyl acetate (99:1) to give protected ketone **17** (80 mg, 84%) as a colorless oil. **17**: $[\alpha]_{\text{D}}^{24} -12$ (*c* 1.4, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.076 [9H, s, Si(CH₃)₃], 0.75 (3H, s, 7a-CH₃), 0.87 (6H, d, *J* = 6.6 Hz, 6'-CH₃ and 7'-H₃), 1.32 (3H, s, 1'-H₃), 2.39 (1H, dd, *J* = 10.8, 6.9 Hz, 3a-H); ¹³C NMR (75 MHz) δ 2.8, 14.4,

18.8, 21.8, 22.6, 22.7, 23.2, 24.0, 27.8, 27.9, 39.6, 39.8, 40.9, 45.0, 49.9, 57.4, 62.3, 78.0, 212.3; HRMS (ESI) calcd. for C₂₁H₄₀O₂SiNa (M⁺+ Na) 375.2695, measured 375.2702.

(1S,3aR,7aR)-7a-Methyl-1-[(S)-6'-methyl-6'-[(triethylsilyl)oxy]-2'-[(trimethylsilyl)oxy]heptan-2'-yl]octahydro-4H-inden-4-one (18).—Protection

of tertiary hydroxyl group in **16** (115 mg, 0.281 mmol) was performed analogously as described above for **15**. The crude product was purified by column chromatography over silica using hexane/ethyl acetate (99:1) to give protected ketone **18** (111 mg, 82%) as a colorless oil. **18**: [α]_D²⁴−9 (*c* 1.0, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.078 [9H, s, Si(CH₃)₃], 0.57 (6H, q, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 0.75 (3H, s, 7a-CH₃), 0.95 (9H, t, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 1.20 (6H, s, 6'-CH₃ and 7'-H₃), 1.32 (3H, s, 1'-H₃), 2.38 (1H, dd, *J* = 10.8, 6.8 Hz, 3a-H); ¹³C NMR (75 MHz) δ 2.8, 6.8, 7.1, 14.4, 18.8, 20.1, 21.9, 24.0, 27.8, 29.9, 30.1, 39.6, 40.9, 45.2, 45.8, 49.9, 57.5, 62.4, 73.2, 78.1, 212.4; HRMS (ESI) calcd. for C₂₇H₅₄O₃Si₂Na (M⁺+ Na) 505.3509, measured 505.3515.

(20S)-3-[(tert-Butyldimethylsilyl)oxy]-20-[(trimethylsilyl)oxy]vitamin D₃ (19).—

To a solution of phosphine oxide **9** (395 mg, 0.860 mmol) in anhydrous THF (5 mL) was added *n*-BuLi (540 μ L, 0.870 mmol) at −78 °C and the stirring of a deep red solution was continued for 30 min. A solution of the ketone **17** (154 mg, 0.430 mmol) in anhydrous THF (2 mL) was then added and the stirring was continued for 3 h. Saturated NH₄Cl (3 mL) was added and the mixture was extracted with diethyl ether (3 × 5 mL). The combined organic layers were washed with water (5 mL), dried (MgSO₄) and evaporated. The crude product was purified by column chromatography using hexane to give protected vitamin D compound **19** (204 mg, 81%) as a colorless oil. **19**: [α]_D²⁴+61 (*c* 1.5, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.070 [9H, s, Si(CH₃)₃], 0.67 (3H, s, 18-H₃), 0.88 (6H, d, *J* = 6.6 Hz, 26- and 27-H₃), 0.90 (9H, s, Si-*t*Bu), 1.29 (3H, s, 21-H₃), 2.12 (1H, m, 1 α -H), 2.26 (1H, dd, *J* = 13.2, 9.6 Hz, 4 β -H), 2.36 (1H, dt, *J* = 13.5, 4.6 Hz, 1 β -H), 2.46 (1H, dd, *J* = 13.2, 4.5 Hz, 4 α -H), 2.82 (1H, br d, *J* ~ 13 Hz, 9 β -H), 3.82 (1H, tt, *J* = 9.6, 4.5 Hz, 3 α -H), 4.80 and 5.01 (1H and 1H, each s, 19-H₂), 6.01 and 6.17 (1H and 1H, each d, *J* = 11.2 Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ −4.6, −4.5, 2.8, 13.9, 18.2, 21.9, 22.0, 22.6, 22.8, 23.3, 23.4, 25.9, 27.7, 27.9, 28.8, 32.8, 36.5, 39.8, 41.2, 45.2, 45.9, 46.9, 56.8, 57.5, 70.7, 78.5, 112.2, 118.1, 121.4, 136.2, 141.5, 145.4; HRMS (ESI) calcd. for C₃₆H₆₆O₂Si₂Na (M⁺+ Na) 609.4499, measured 609.4508.

(20S)-3-[(tert-Butyldimethylsilyl)oxy]-25-[(triethylsilyl)oxy]-20-[(trimethylsilyl)oxy]-vitamin D₃ (20).—The Wittig-Horner reaction between

ketone **18** (44 mg, 0.092 mmol) and phosphine oxide **9** (83 mg, 0.184 mmol) was performed analogously as described above for **17**. The crude product was purified by column chromatography over silica using hexane/ethyl acetate (99:1) to give protected vitamin D compound **20** (60 mg, 91%) as a colorless oil. **20**: [α]_D²⁴+54 (*c* 1.0, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.069 [9H, s, Si(CH₃)₃], 0.57 (6H, q, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 0.66 (3H, s, 18-H₃), 0.90 (9H, s, Si-*t*Bu), 0.95 (9H, t, *J* = 7.5 Hz, 3 × SiCH₂CH₃), 1.20 (6H, s, 26- and 27-H₃), 1.30 (3H, s, 21-H₃), 2.12 (1H, m, 1 α -H), 2.23 (1H, dd, *J* = 13.2, 9.3 Hz, 4 β -H), 2.36 (1H, dt, *J* = 13.7, 4.8 Hz, 1 β -H), 2.46 (1H, dd, *J* = 13.2, 4.2 Hz, 4 α -H), 2.82 (1H, br d, *J* ~ 12

Hz, 9 β -H), 3.81 (1H, tt, J = 9.3, 4.2 Hz, 3 α -H), 4.80 and 5.00 (1H and 1H, each s, 19-H₂), 6.01 and 6.17 (1H and 1H, each d, J = 11.2 Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ -4.6, -4.5, 2.8, 6.8, 7.2, 13.9, 18.2, 20.2, 21.9, 22.0, 23.4, 25.9, 26.9, 27.7, 28.9, 29.8, 30.1, 32.8, 36.4, 41.1, 45.3, 45.8, 46.9, 56.8, 57.5, 70.7, 73.3, 78.6, 112.1, 118.1, 121.4, 136.2, 141.6, 145.4; HRMS (ESI) calcd. for C₄₂H₈₀O₃Si₃Na (M⁺+ Na) 739.5313, measured 739.5322.

(20S)-20-Hydroxyvitamin D₃ (3).—Tetra-*n*-butylammonium fluoride (1M in THF; 4.3 mL, 4.3 mmol) was added to a solution of protected vitamin D compound **23** (84 mg, 0.143 mmol) in anhydrous THF (5 mL) and the mixture was stirred at room temperature overnight. Brine (3 mL) was added and the mixture was extracted with AcOEt (3 \times 10 mL). The combined organic layers were washed with water (5 mL), dried (MgSO₄) and evaporated. The crude product was purified by HPLC (9.4 mm \times 25 cm Zorbax-Sil column, 4 mL/min) using hexane/2-propanol (95:5) solvent system; vitamin **3** (48 mg, 84%) was collected at R_V = 23 mL. Analytical sample of the vitamin was obtained after reversed-phase HPLC (9.4 mm \times 25 cm Zorbax Eclipse XDB-C18 column, 4 mL/min) using methanol/water (9:1) solvent system (R_V = 44 mL). **3**: [α]_D²⁴+21 (*c* 1.1, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.72 (3H, s, 18-H₃), 0.87 (6H, d, J = 6.6 Hz, 26- and 27-H₃), 1.27 (3H, s, 21-H₃), 2.17 (1H, ddd, J = 13.5, 8.5, 4.8 Hz, 1 α -H), 2.28 (1H, dd, J = 13.2, 7.2 Hz, 4 β -H), 2.39 (1H, ddd, J = 13.5, 7.6, 4.8 Hz, 1 β -H), 2.57 (1H, dd, J = 13.2, 3.6 Hz, 4 α -H), 2.82 (1H, br d, J ~ 13 Hz, 9 β -H), 3.94 (1H, tt, J = 7.3, 3.6 Hz, 3 α -H), 4.82 and 5.04 (1H and 1H, each s, 19-H₂), 6.03 and 6.23 (1H and 1H, each d, J = 11.2 Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ 13.7, 21.9, 22.0, 22.1, 22.6, 22.7, 23.4, 26.6, 27.9, 28.9, 31.9, 35.2, 39.6, 40.8, 43.9, 45.9, 46.0, 56.5, 58.2, 69.2, 75.2, 112.5, 118.2, 122.3, 135.4, 141.5, 145.0; HRMS (ESI) calcd. for C₂₇H₄₅O₂ (M⁺+ Na) 401.3420, measured 401.3428.

(20S)-20,25-Dihydroxyvitamin D₃ (6).—The hydroxyl deprotection in the vitamin D compound **24** (60 mg, 0.083 mmol) was performed analogously as described above for **23**. The crude product was purified by HPLC (9.4 mm \times 25 cm Zorbax-Sil column, 4 mL/min) using hexane/2-propanol (9:1) solvent system; vitamin **6** (26 mg, 75%) was collected at R_V = 29 mL. Analytical sample of the vitamin was obtained after reversed-phase HPLC (9.4 mm \times 25 cm Zorbax Eclipse XDB-C18 column, 4 mL/min) using methanol/water (8:2) solvent system (R_V = 37 mL). **6**: [α]_D²⁴+20 (*c* 1.3, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.72 (3H, s, 18-H₃), 1.22 (6H, s, 26- and 27-H₃), 1.28 (3H, s, 21-H₃), 2.17 (1H, ddd, J = 13.4, 8.5, 4.6 Hz, 1 α -H), 2.28 (1H, dd, J = 13.2, 7.3 Hz, 4 β -H), 2.40 (1H, ddd, J = 13.4, 7.5, 4.6 Hz, 1 β -H), 2.57 (1H, dd, J = 13.2, 3.5 Hz, 4 α -H), 2.81 (1H, br d, J ~ 13 Hz, 9 β -H), 3.94 (1H, tt, J = 7.3, 3.5 Hz, 3 α -H), 4.81 and 5.05 (1H and 1H, each s, 19-H₂), 6.03 and 6.22 (1H and 1H, each d, J = 11.2 Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ 13.8, 18.9, 21.9, 22.0, 23.3, 26.5, 28.8, 29.2, 29.5, 31.9, 35.2, 40.8, 43.9, 44.4, 45.8, 46.0, 56.5, 58.3, 69.2, 71.0, 75.2, 112.5, 118.2, 122.3, 135.5, 141.4, 145.1; HRMS (ESI) calcd. for C₂₇H₄₅O₃ (M⁺+ H) 417.3369, measured 417.3372.

(3S)-3-[(1'S,3a'R,4'S,7a'S)-7a'-Methyl-4'-[(triethylsilyl)oxy]-octahydro-1H-inden-1'-yl]-3-hydroxybutanenitrile (21).—To a solution of acetonitrile (0.45 mL, 8.69 mmol) in anhydrous THF (20 mL) was added LDA (2M in THF/heptane/ethylbenzene; 4.3 mL, 8.60 mmol) at -78 °C. The yellow

mixture was stirred at this temperature for 30 min. A solution of ketone **11** (0.60 g, 1.93 mmol) in anhydrous THF (10 mL) was added and stirring was continued for 2 h. Saturated NH_4Cl (10 mL) was slowly added and the mixture was extracted with ethyl acetate (3×15 mL). The combined organic layers were washed with water (10 mL), dried (MgSO_4) and evaporated. The residue was purified by column chromatography on silica using hexane/ethyl acetate (85:15) to give hydroxy nitrile **21** (0.624 g, 92%) as a colorless oil. **21**: $[\alpha]_{\text{D}}^{24} +29$ (c 1.0, CHCl_3); ^1H NMR (300 MHz, CDCl_3) δ 0.54 (6H, q, $J = 7.8$ Hz, SiCH_2CH_3), 0.94 (9H, t, $J = 7.8$ Hz, SiCH_2CH_3), 1.11 (3H, s, $7\text{a}'\text{-CH}_3$), 1.49 (3H, s, 4-H_3), 2.43 and 2.49 (1H and 1H, each d, $J = 16.5$ Hz, 2-H_2), 4.05 (1H, narr m, $4'\alpha\text{-H}$); ^{13}C NMR (75 MHz) δ 4.8, 6.9, 15.4, 17.5, 21.6, 22.6, 27.1, 32.4, 34.3, 40.8, 42.9, 52.9, 57.8, 69.2, 73.6, 117.9; HRMS (ESI) calcd. for $\text{C}_{20}\text{H}_{37}\text{NO}_2\text{SiNa}$ ($\text{M}^+ + \text{Na}$) 374.2491, measured 374.2495.

(3S)-3-[(1'S,3a'R,4'S,7a'S)-4'-Hydroxy-7a'-methyl-octahydro-1H-inden-1'-yl)]-3-hydroxybutanenitrile (22).—

To a solution of nitrile **21** (0.458 g, 1.30 mmol) in CH_2Cl_2 (20 mL) was added a solution of hydrogen chloride in dioxane (4 M; 0.98 mL, 3.92 mmol) at 0°C and the mixture was stirred for 30 min. Then, water (5 mL) and NaHCO_3 (10 mL) were added and the layers were separated. The aqueous phase was extracted with dichloromethane (2×10 mL) and the combined organic layers were washed with water (10 mL), dried (MgSO_4) and evaporated. The residue was purified by column chromatography on silica using hexane/ethyl acetate (7:3) to give diol **22** (0.302 g, 98%) as a colorless oil. **22**: $[\alpha]_{\text{D}}^{24} +14$ (c 1.0, CHCl_3); ^1H NMR (300 MHz, CDCl_3) δ 1.13 (3H, s, $7\text{a}'\text{-CH}_3$), 1.51 (3H, s, 4-H_3), 2.44 and 2.50 (1H and 1H, each d, $J = 16.6$ Hz, 2-H_2), 4.10 (1H, narr m, $4'\text{-H}$); ^{13}C NMR (75 MHz) δ 15.3, 17.3, 21.5, 22.1, 27.1, 32.6, 33.4, 40.5, 42.8, 52.4, 57.6, 69.1, 73.5, 117.8; HRMS (ESI) calcd. for $\text{C}_{14}\text{H}_{23}\text{NO}_2\text{Na}$ ($\text{M}^+ + \text{Na}$) 260.1626, measured 260.1620.

(3S)-3-Hydroxy-3-[(1'S,3a'R,7a'R)-7a'-methyl-4'-oxo-octahydro-1H-inden-1'-yl]butanenitrile (23).—

The oxidation of the secondary hydroxyl group in **22** (73 mg, 0.307 mmol) was performed analogously as described above for **13**. The crude product was purified by column chromatography over silica using hexane/ethyl acetate (7:3) to give the ketone **23** (68 mg, 95%) as a colorless oil. **23**: $[\alpha]_{\text{D}}^{24} -19$ (c 0.97, CHCl_3); ^1H NMR (300 MHz, CDCl_3) δ 0.80 (3H, s, $7\text{a}'\text{-CH}_3$), 1.55 (3H, s, 4-H_3), 2.48 and 2.55 (1H and 1H, each d, $J = 16.5$ Hz, 2-H_2); ^{13}C NMR (75 MHz) δ 14.2, 18.8, 21.9, 23.7, 27.5, 32.5, 38.9, 40.7, 49.7, 57.7, 61.8, 73.0, 117.6, 211.1; HRMS (ESI) calcd. for $\text{C}_{14}\text{H}_{21}\text{NO}_2\text{Na}$ ($\text{M}^+ + \text{Na}$) 258.1470, measured 258.1465.

(3S)-3-[(1'S,3a'R,7a'R)-7a'-methyl-4'-oxo-octahydro-1H-inden-1'-yl)]-3-[(trimethylsilyloxy)butanenitrile (24).—

To a solution of hydroxy nitrile **23** (89 mg, 0.378 mmol) in anhydrous CH_2Cl_2 (4 mL) was added pyridine (300 μL , 3.78 mmol) and trimethylsilyl trifluoromethanesulfonate (206 μL , 1.14 mmol) at 0°C and the mixture was stirred at this temperature for 1 h. Saturated NaHCO_3 (3 mL) was added and the mixture was extracted with dichloromethane (3×5 mL). The combined organic layers were washed with 5% HCl (5 mL), water (5 mL), dried (MgSO_4) and evaporated. The residue was purified by column chromatography on silica using hexane/ethyl acetate (9:1) to give ketone **24** (101 mg, 86%) as a colorless oil. **24**: $[\alpha]_{\text{D}}^{24} -15$

(*c* 0.97, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.14 [9H, s, Si(CH₃)₃], 0.75 (3H, s, 7a'-CH₃), 1.57 (3H, s, 4-H₃), 2.41 and 2.49 (1H and 1H, each d, *J* = 16.4 Hz, 2-H₂); ¹³C NMR (75 MHz) δ 2.4, 14.2, 18.8, 22.0, 23.8, 27.9, 32.5, 39.2, 40.7, 49.7, 58.5, 61.9, 75.9, 117.8, 211.4; HRMS (ESI) calcd. for C₁₇H₂₉NO₂SiNa (M⁺ + Na) 330.1865, measured 330.1870.

(20S)-3-[(*tert*-Butyldimethylsilyloxy]-22-cyano-20-[(trimethylsilyloxy]-23,24,25,26,27-pentanorvitamin D₃

(25).—The Wittig-Horner reaction between ketone **24** (100 mg, 0.325 mmol) and phosphine oxide **11** (294 mg, 0.650 mmol) was performed analogously as described above for **17**. The crude product was purified by column chromatography over silica using hexane/ethyl acetate (96:4) to give protected vitamin D compound **25** (152 mg, 87%) as a colorless oil. **25**: [α]_D²⁴ +52.6 (*c* 1.0, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.064 and 0.070 (3H and 3H, each s, 2 × SiCH₃), 0.14 [9H, s, Si(CH₃)₃], 0.66 (3H, s, 18-H₃), 0.88 (9H, s, Si-*t*-Bu), 1.54 (3H, s, 21-H₃), 2.23 (1H, dd, *J* = 13.0, 9.1 Hz, 4β-H), 2.36 (1H, m, 1β-H), 2.40 and 2.55 (1H and 1H, each d, *J* = 16.3 Hz, 22-H₂), 2.44 (1H, dd, *J* = 13.0, 3.1 Hz, 4α-H), 2.82 (1H, br d, *J* ~ 12 Hz, 9β-H), 3.82 (1H, tt, *J* = 9.1, 4.3 Hz, 3α-H), 4.78 and 5.01 (1H and 1H, each s, 19-H₂), 6.01 and 6.15 (1H, and 1H, each d, *J* = 11.2 Hz, 6- and 7-H); ¹³C NMR (75 MHz) δ -4.6, -4.5, 2.5, 13.7, 18.2, 21.9, 22.3, 23.2, 25.9, 27.8, 28.7, 32.5, 32.8, 36.4, 40.8, 45.9, 46.9, 56.4, 58.9, 70.6, 76.5, 112.2, 118.2, 118.6, 121.2, 136.8, 140.5, 145.5; HRMS (ESI) calcd. for C₃₂H₅₅NO₂Si₂Na (M⁺ + Na) 564.3669, measured 564.3675.

(20S)-3-[(*tert*-Butyldimethylsilyloxy]-22-formyl-20-[(trimethylsilyloxy]-23,24,25,26,27-pentanorvitamin D₃

(26).—To a solution of nitrile **25** (105 mg, 0.193 mmol) in anhydrous CH₂Cl₂ (5 mL) was added diisobutylaluminum hydride (1.20 mL, 1.20 mmol) at -10 °C. Potassium sodium tartrate solution (2 mL) was added after 1 h and the mixture was extracted with CH₂Cl₂ (3 × 5 mL). The combined organic layers were washed with water, dried (MgSO₄) and evaporated. The crude product was purified by column chromatography using hexane/ethyl acetate (98:2) to give aldehyde **26** (65 mg, 62%) as a colorless oil. **26**: [α]_D²⁴ +50.5 (*c* 1.4, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.08 and 0.09 (3H and 3H, each s, 2 × SiCH₃), 0.14 [9H, s, Si(CH₃)₃], 0.70 (3H, s, 18-H₃), 0.91 (9H, s, Si-*t*-Bu), 1.53 (3H, s, 21-H₃), 2.25 (1H, m, 4β-H), 2.39 (1H, dt, *J* = 13.6, 4.8 Hz, 1β-H), 2.42 (1H, dd, *J* = 13.2, 4.5 Hz, 4α-H), 2.50 and 2.65 (1H and 1H, each dd, *J* = 14.0, 3.4 Hz, 22-H₂), 2.83 (1H, br d, *J* ~ 11.5 Hz, 9β-H), 3.84 (1H, tt, *J* = 9.0, 4.0 Hz, 3α-H), 4.80 and 5.03 (1H and 1H, each m, 19-H₂), 6.03 and 6.17 (1H and 1H, each d, *J* = 11.2 Hz, 7- and 6-H), 9.79 (1H, t, *J* = 3.4 Hz, CHO); ¹³C NMR (75 MHz) δ -4.61, -4.60, 2.6, 14.0, 18.2, 21.9, 22.6, 23.3, 25.9, 28.7, 28.8, 32.8, 36.4, 41.0, 46.1, 46.9, 56.5, 57.5, 60.4, 70.6, 76.9, 112.2, 118.5, 121.2, 136.6, 140.7, 145.4, 202.1; HRMS (ESI) calcd. for C₃₂H₅₆O₃Si₂Na (M⁺ + Na) 567.3666, measured 567.3678.

(20S,23R)- and (20S,23S)-3-[(*tert*-Butyldimethylsilyloxy]-23-hydroxy-20-[(trimethylsilyloxy]-vitamin D₃ (27a,b).

—To a solution of aldehyde **26** (37 mg, 0.068 mmol) in anhydrous THF (2 mL) was added isobutylmagnesium chloride (2 M in THF; 68 μL, 0.136 mmol) at 0 °C and the mixture was stirred at rt overnight. Then, saturated NH₄Cl (1 mL) was added and the mixture was extracted with AcOEt (3 × 3 mL). The combined organic layers were washed with water (2 mL), dried (MgSO₄) and evaporated. The crude

product was purified by column chromatography using hexane/ethyl acetate (95:5) to give mixture of alcohols **27a,b** (29 mg, 70%) as a colorless oil.

(20S,23R)- and (20S,23S)-20,23-Dihydroxyvitamin D₃ (4a,b).—The hydroxyl deprotection in the vitamin D compounds **31a,b** (54 mg, 0.090 mmol) was performed analogously as described above for **23**. The products were separated by RP-HPLC (9.4 mm × 25 cm Zorbax Eclipse XDB-C18 column, 4 mL/min) using methanol/water (85:15) solvent system. Pure vitamin **4b** (16 mg) was collected at $R_V = 28$ mL and **4a** (11 mg) at $R_V = 48$ mL (total yield 74%).

4a: $[\alpha]_D^{24} +25.1$ (c 0.9, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.63 (3H, s, 18-H₃), 0.92 (6H, d, $J = 6.6$ Hz, 26- and 27-H₃), 1.30 (3H, s, 21-H₃), 2.17 (1H, ddd, $J = 13.1, 8.6, 4.7$ Hz, 1α-H), 2.27 (1H, dd, $J = 13.1, 7.6$ Hz, 4β-H), 2.39 (1H, m, 1β-H), 2.56 (1H, dd, $J = 13.1, 3.8$ Hz, 4α-H), 2.82 (1H, dd, $J = 11.8, 3.9$ Hz, 9β-H), 3.93 (1H, m, 3α-H), 4.13 (1H, m, 23-H), 4.80 and 5.04 (1H and 1H, each m, 19-H₂), 6.03 and 6.21 (1H and 1H, each d, $J = 11.2$ Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ 14.1, 21.9, 22.1, 22.5, 23.3, 23.6, 24.2, 28.2, 28.8, 31.9, 35.2, 40.7, 45.6, 45.9, 47.9, 49.2, 56.5, 67.2, 69.2, 76.2, 112.5, 118.4, 122.2, 135.7, 141.2, 145.1; HRMS (ESI) calcd. for C₂₇H₄₅O₃ (M⁺ + H) 417.3369, measured 417.3374.

4b: $[\alpha]_D^{24} +17.6$ (c 0.9, CHCl₃); ¹H NMR (300 MHz, CDCl₃) δ 0.71 (3H, s, 18-H₃), 0.90 and 0.92 (6H, each d, $J = 6.6$ Hz, 26- and 27-H₃), 1.39 (3H, s, 21-H₃), 2.19 (1H, ddd, $J = 13.1, 8.7, 4.8$ Hz, 1α-H), 2.27 (1H, dd, $J = 13.1, 7.5$ Hz, 4β-H), 2.39 (1H, m, 1β-H), 2.56 (1H, dd, $J = 13.1, 3.8$ Hz, 4α-H), 2.81 (1H, br d, $J \sim 12$ Hz, 9β-H), 3.93 (1H, m, 3α-H), 4.09 (1H, m, 23-H), 4.80 and 5.04 (1H and 1H, each m, 19-H₂), 6.03 and 6.21 (1H and 1H, each d, $J = 11.2$ Hz, 7- and 6-H); ¹³C NMR (75 MHz) δ 13.8, 21.9, 22.2, 22.4, 23.2, 23.3, 24.3, 26.0, 28.8, 31.9, 35.2, 40.9, 45.9, 46.0, 46.9, 47.4, 56.5, 61.2, 66.8, 69.2, 76.8, 112.5, 118.3, 122.2, 135.6, 141.2, 145.1; HRMS (ESI) calcd. for C₂₇H₄₅O₃ (M⁺ + H) 417.3369, measured 417.3375.

Measurement of CYP27B1 activity on secosteroids

Mouse CYP27B1 was expressed in *E. coli* and purified as before [68]. CYP27B1 activity was measured on substrate incorporated into the membrane of phospholipid vesicles as described previously [68]. Briefly, phospholipid vesicles were prepared by sonication of dioleoyl phosphatidylcholine, cardiolipin and secosteroid. Vesicles (510 μM phospholipid) were then incubated at 37°C with CYP27B1 (0.2 μM) in buffer comprising 20 mM Hepes, pH 7.4, 100 mM NaCl, 0.1 mM EDTA, 0.1 mM and dithiothreitol, adrenodoxin (15 μM), adrenodoxin reductase (0.4 μM), glucose-6-phosphate (2 mM), glucose-6-phosphate dehydrogenase (2 U/mL) and NADPH (50 μM) in a volume of 0.5 ml. After reaction (5 or 20 min), the incubation mixture was extracted 3-times with 2.5 ml dichloromethane, extracts combined, dried under N₂ gas and dissolved in 64% methanol. Samples were analysed by reverse phase HPLC on a Grace Alltima column (25 × 0.46 cm) using a gradient of 64 to 100% methanol for 15 min followed by 100% methanol for 30 min, at a flow rate of 0.5 ml/min. Metabolites were detected with a UV detector at 265 nm.

Nuclear receptor analyses

VDR translocation assay—To study the ligand-induced VDR translocation from cytoplasm to the nucleus, we used SKMEL-188 melanoma cells transduced with pLenti-CMV-VDR-EGFP-pgk-puro, which express VDR-GFP fusion protein, and followed the protocols described previously [57, 80, 81]. Briefly, cells were grown on a 96 well plate using Ham's F10 media containing 5% charcoal-treated FBS until 70% confluence. The cells were treated with the secosteroids for 1.5 h in a 96 well plate followed by treatment of NucBlue™ Live ReadyProbes™ Reagent (Invitrogen, Waltham, MA, USA) using the manufacturer's protocol. The ratio of cell number with a fluorescent nucleus (GFP) to the total cell number (DAPI) was determined using Cytation 5 (Biotek, Winooski, VT, USA).

AhR binding assay—The binding of chemically synthesized secosteroids to AhR was tested using the Human AhR Reporter Assay System (Indigo Biosciences, State College, PA, USA) following the manufacturer's protocol and as detailed previously [62]. Briefly, the cells including the luciferase reporter gene linked to an AhR-responsive promoter were treated with the compounds for 23 h and the luminescence signal was measured using Cytation 5 (Biotek, Winooski, VT, USA) after treatment with the detection substrate provided in the kit system.

LXR binding assay—The bindings were performed using the LanthaScreen TR-FRET LXR α / β Coactivator kit (Thermo Fisger Scientific, Inc., Waltham, MA) as described previously [61]. Briefly, TR-FRET ratios were calculated by dividing the emission at 520 nm by that at 495 nm using Synergy neo2 (BioTek, Winooski, VT).

Molecular modeling—To predict binding poses of vitamin D₃ derivatives: 20*S*(OH)D₃, 20*S*,23*S*(OH)₂D₃, 20*S*,23*R*(OH)₂D₃ and 20*S*,25(OH)₂D₃ towards various receptors including VDR, ROR α and ROR γ , liver X receptors (LXR α and LXR β) and aryl hydrocarbon receptor (AhR), a receptor-based approach (molecular docking) was used. Crystal structures for VDR (PDBID: 1DB1) [82], ROR α (PDBID: 1S0X) [83], ROR γ (PDBID: 3L0J) [84], LXR α (PDBID: 5AVI) [85] and LXR β (PDBID: 5HJP) [86], and a human AhR homology model from our previous study [62] were used for the docking studies. Openbabel [87] software was used to convert the ligands of vitamin D₃ derivatives in PDB format to PDBQT format. Receptors were loaded into AutoDock Tools [88] software for preparation: polar hydrogen atoms were added; water molecules, ion molecules, and bound ATP were removed; and lastly, Kollman Charges were added. Prepared receptors were output in PDBQT format for molecular docking in Autodock Vina [89]. To evaluate the receptor-ligand interaction, 3D and 2D interactions between vitamin D₃ derivatives with each studied receptor were performed using PyMol [90] software for 3D interactions and Maestro [91] for 2D interaction map.

Cell culture studies

Proliferation—Human dermal fibroblasts isolated from the skin of black neonates (HDFn) and human epidermal (HaCaT) keratinocytes cultured as described previously [50, 80] were used for the experiments. Briefly, HDFn were plated onto 96-well plates at a confluence of about 70% in their seventh passage. Both HDFn and HaCaT keratinocytes were treated

with $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$ or $1\alpha,25(OH)_2D_3$ (Sigma-Aldrich, St. Louis, MO, USA) dissolved in ethanol and diluted in DMEM containing charcoal-treated serum, as previously described [44, 45, 80]. The vehicle control comprised 0.1% ethanol. Proliferation was estimated using MTS (Promega, Madison, WI) assay [44, 45, 80]. The number of viable cells was measured in 6 replicates.

Gene expression—RNA isolation and reverse transcription were performed as described previously [61]. Quantitative RT-PCR was carried out using Sybr green Master Mix (Thermo Scientific, Waltham, MA), in triplicates as previously described [61, 92]. Human epidermal keratinocytes (HEKn) isolated from foreskin of white neonates were cultured until the third passage before treatment as described in the literature [46, 70]. Cells were treated with vitamin D₃ derivatives for 8 h or 24 h and harvested for RNA isolation. Approximately 300 ng of RNA was used for cDNA synthesis and an equal amount of cDNA was used for PCR reactions. Cyclophilin B or GAPDH was used as an internal control. Primers sequences are listed in [45, 81]. Reactions were done in triplicate.

Protein expression—HEKn were cultured until the third passage before treatment. Keratinocytes were plated in a 96-well plate and at 80% confluence, 10^{-7} M secosteroids (or 0.1% ethanol control) were added and cells were incubated for 48 h. Cells were fixed and stained with anti-involucrin (1:20, GTX-14504; Genetex, Irvine, CA, USA) anti-CK10 (1:300, Santa Cruz Biotechnology), anti-catalase (1:200, Santa Cruz Biotechnology) or anti-CK14 antibody with Alexa Fluor[®] 488 (1:500, Santa Cruz Biotechnology) and imaged at 10× magnification in Cytation 5 reader as described before [45, 81, 93]. Nuclei were stained with blue DAPI. Fluorescence intensity was measured using the Cytation 5 reader and ImageJ, and data were analyzed using Graph Pad Prism.

Human and T cell activation and analysis—Human peripheral blood mononuclear cells (PBMC) from two healthy individuals were obtained from a pre-existing repository. Their use was approved by the University of Alabama at Birmingham Institutional Review Board (Human Subject Assurance Number FWA00005960) as an exempt protocol #4 (Dr C. Raman, P.I.). The protocol was classified for exempt status under 45CFR46.102 (f) in that it does not involve “human subjects” as defined therein. Informed consent is waived in accord with 45CFR46.116(d). In 48 well plates, 0.5×10^6 PBMC/well were cultured in RPMI1640 containing, 5% charcoal adsorbed FBS and antibiotics and T cells were activated using 0.5 µg/ml anti-CD3 (clone 145–2C11) and 1 µg/ml anti-CD28 (clone 37.51) for 24 h [44, 45, 80, 94, 95]. The culture was treated with 10^{-7} M $20S(OH)D_3$, $20S,25(OH)_2D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$ or $1\alpha,25(OH)_2D_3$ (Sigma-Aldrich, St. Louis, MO, USA) dissolved in ethanol and diluted in culture medium at the same time of addition of anti-CD3 and anti-CD28. The vehicle control comprised of 0.1% ethanol. At the end of assay the cells were stained with antibodies to CD4 (clone RPA-T4, BV-421), CD19 (clone HB19, BV-510), CD14 (clone HCD14, Alexa 700), CD137 (clone 4B4–1, BV605), CD8a (clone RPA-T8, FITC), ICOS (clone C398.4A, PE/Dazzle), PD1 (clone NAT105, PE), CD80 (clone 2D10, PerCP/Cy5.5), OX40 (clone ACT35, PE/Cy7), CD69 (clone FN50, APC), HLA-DR (clone L243, APC/Fire 750). All antibodies were from Biolegend except anti-CD8

was from Thermo Fisher. The stained cells were analyzed using a CytoFLEX (Beckman Coulter) flow cytometer.

For unbiased analysis of the multidimensional data, we utilized computational algorithm-based approaches [96]. The flow cytometry standard (FCS) data files from each individual and treatment (two individuals, six treatments including ethanol, a total of 12 data files) were processed through the following workflow – compensation, gating on live cells, gating for single cells and subtraction of CD19⁺ (B cells) and CD14⁺ (monocytes). We removed B cells and monocytes from the analysis because only T cells (CD4⁺ and CD8⁺) are activated by anti-CD3 and anti-CD28. After these initial gating steps, a key word was added to each of the 12 FCS files, down sampled to 50,000 events/sample and concatenated. Down sampling ascertains that each sample is equally weighted, and concatenation ascertains that statistical analysis is applied equally. The concatenated files were first subjected to TriMAP analysis, a computational algorithm that clusters populations based on triplet relatedness followed by FlowSOM algorithm-based clustering [96]. To perform these analyses, we used the plugins within FlowJo (BD, CA, USA).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- more efficient routes of chemical synthesis for $20S(OH)D_3$, $20S,23S(OH)_2D_3$, $20S,23R(OH)_2D_3$ are reported
- new synthetic pathway for $20S,25(OH)_2D_3$ is reported
- phenotypic effects of D3 derivatives are compared
- actions of D3 derivatives on different nuclear receptors are compared
- analysis of D3 derivatives on T cell activation in PBMC is reported

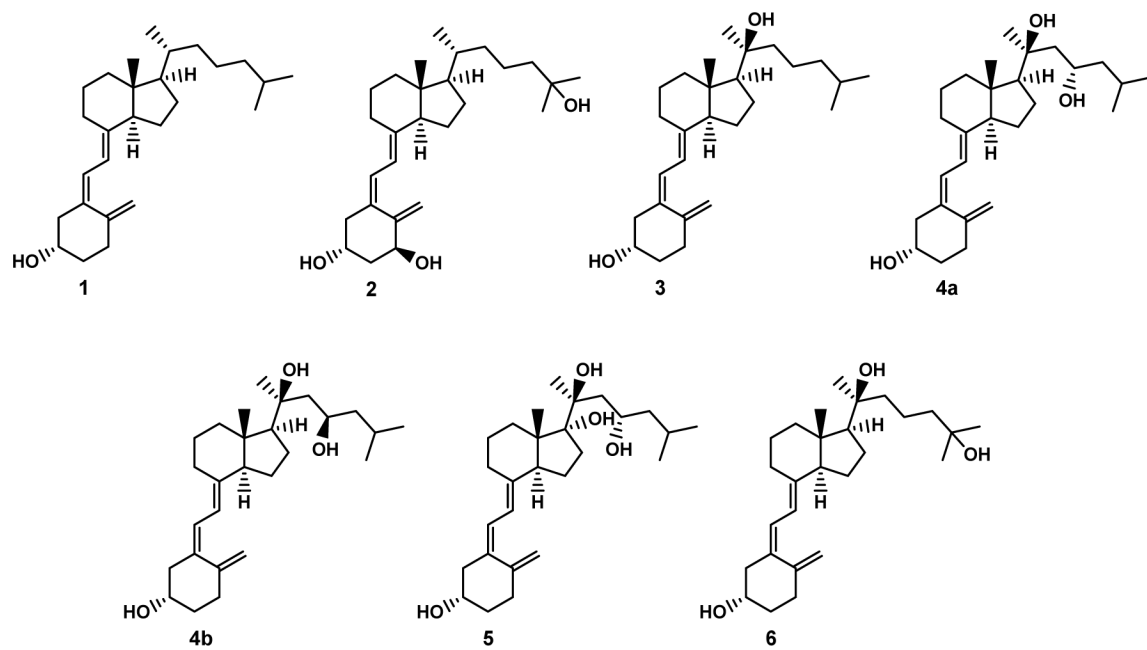


Figure 1.
Chemical structure of vitamin D₃ and some of its biologically important derivatives.

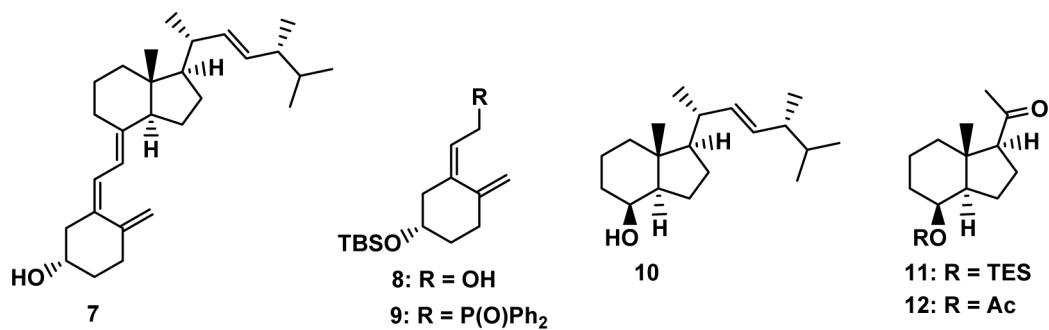


Figure 2.
Vitamin D₂ (7) and the building blocks used as the key intermediates in the synthesis of target vitamin D compounds.

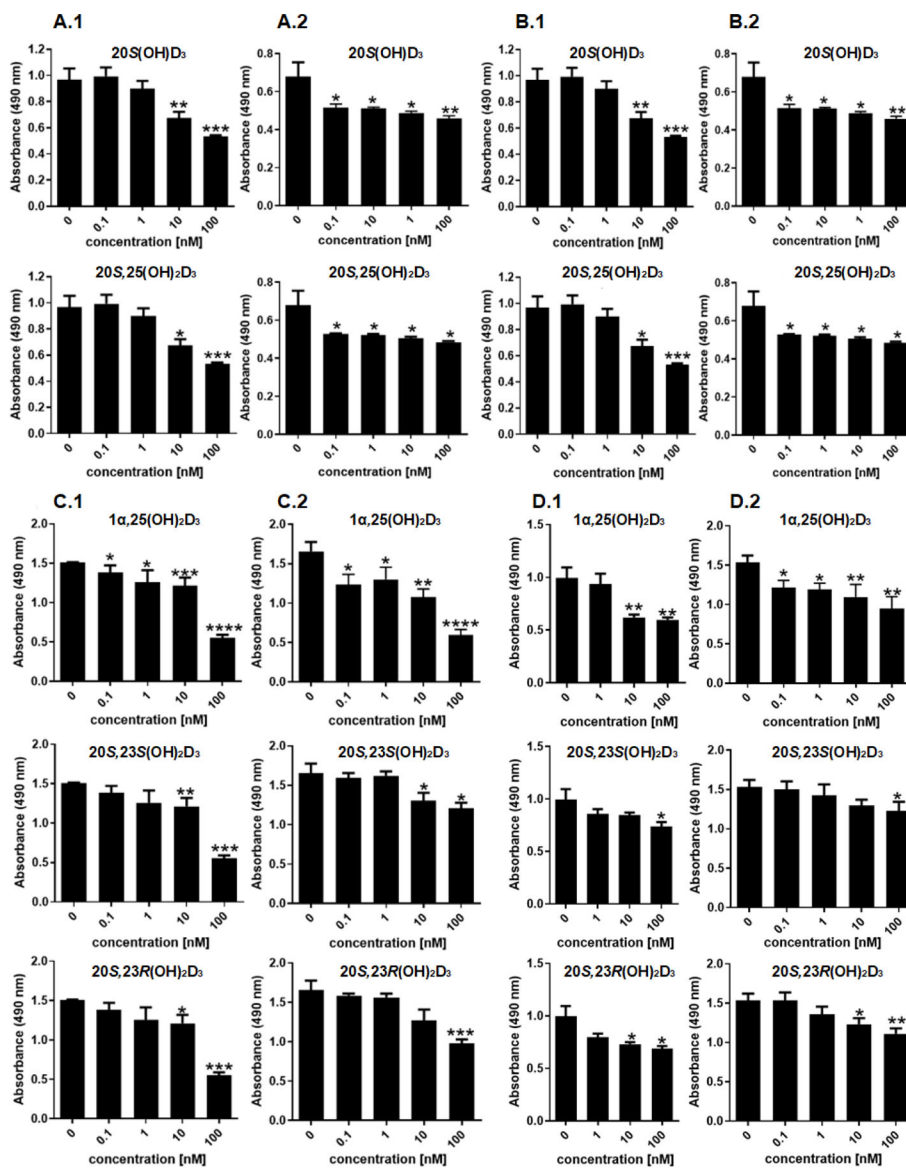


Figure 3. Inhibition of cell proliferation by 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ in comparison to 1α,25(OH)₂D₃. Human epidermal (HaCaT) keratinocytes (A, C) or dermal fibroblasts (B, D) were treated for 24 h (1) or 48 h (2) with graded concentrations of the compounds: 0.1, 1, 10, or 100 nM. Ethanol (0.1%) served as a negative control. Data are means ± SD (n = 6) and were analyzed using Student's *t*-test with *p* < 0.05 (*), *p* < 0.01 (**) or *p* < 0.001 (***).

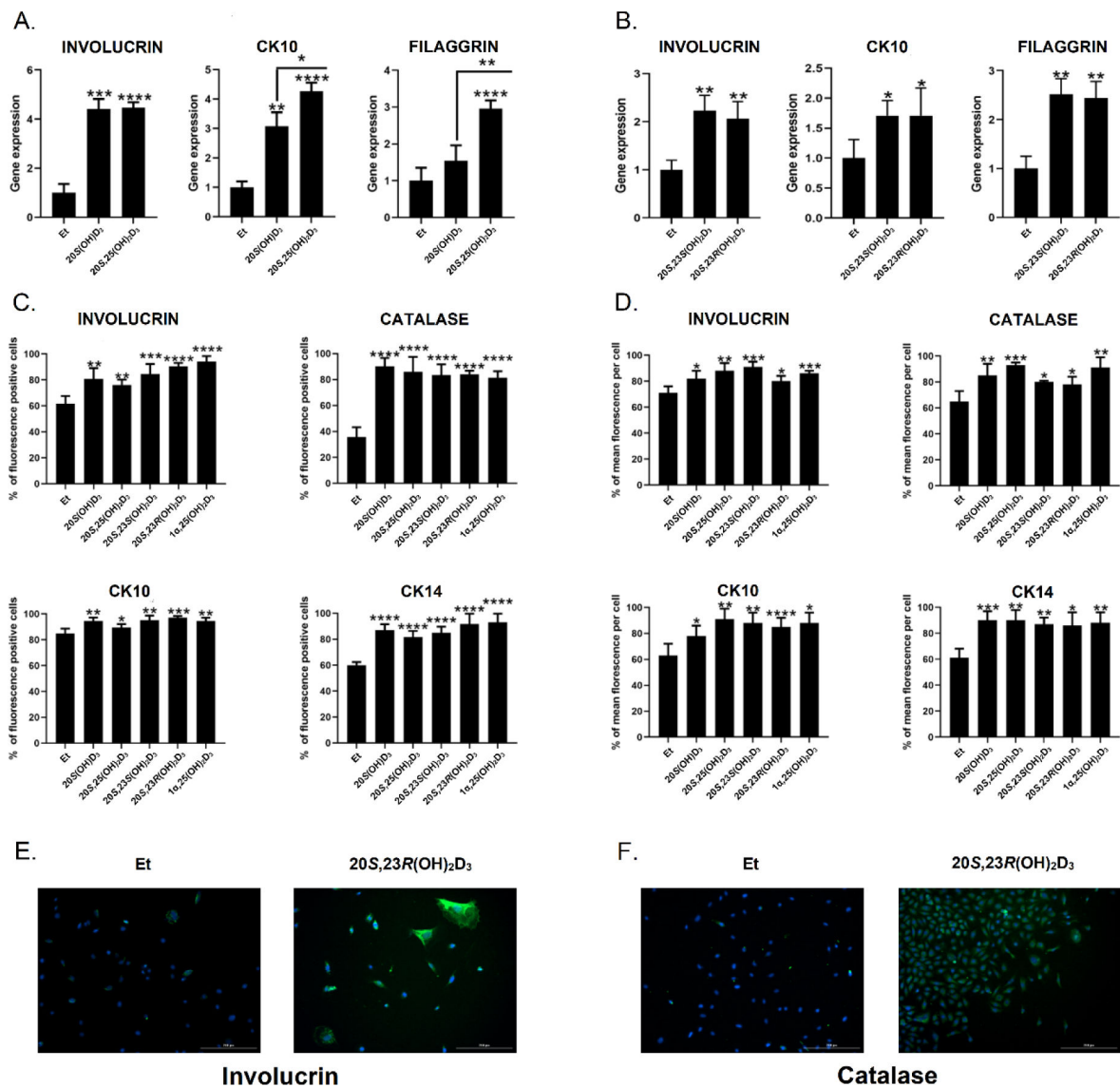
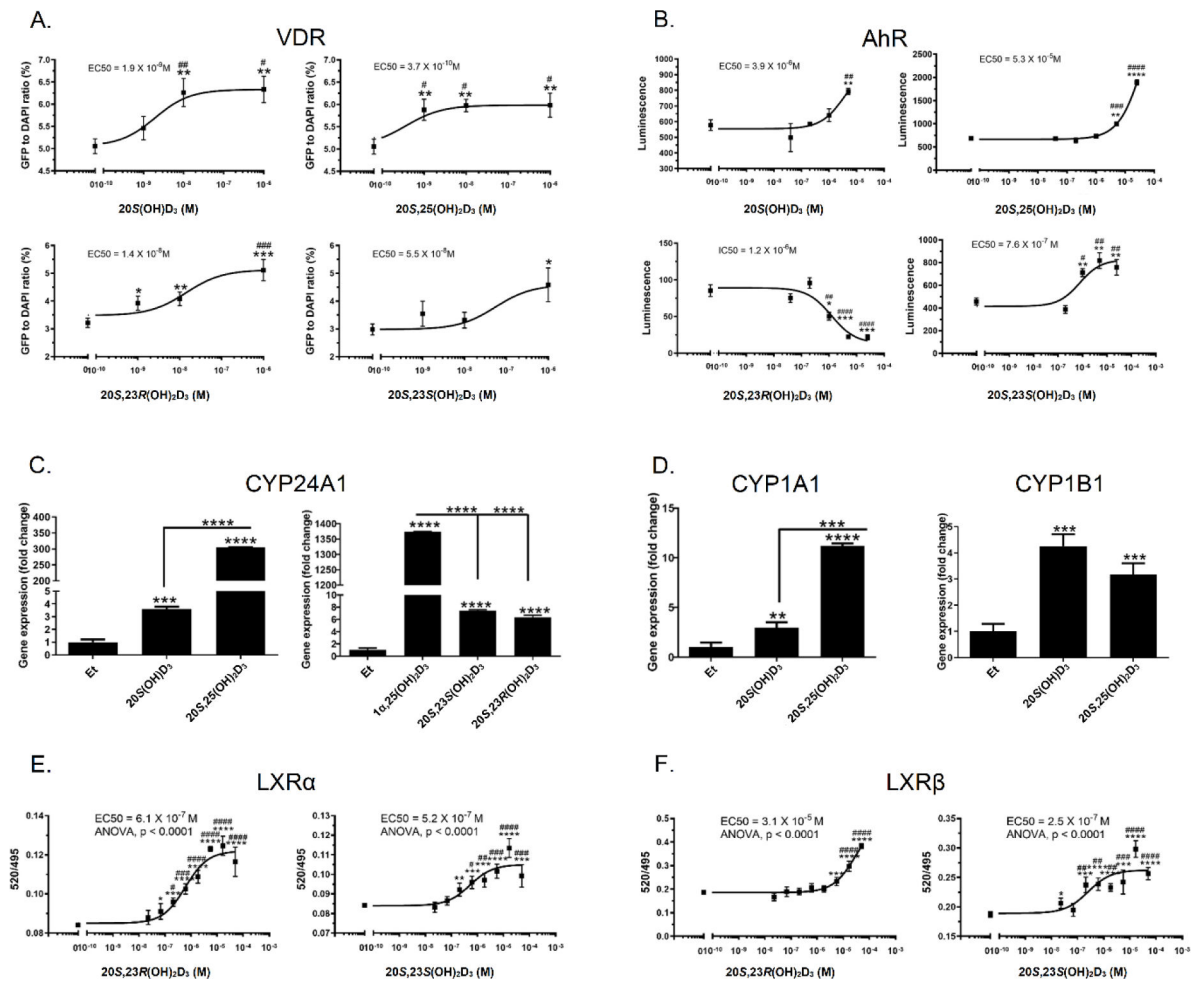
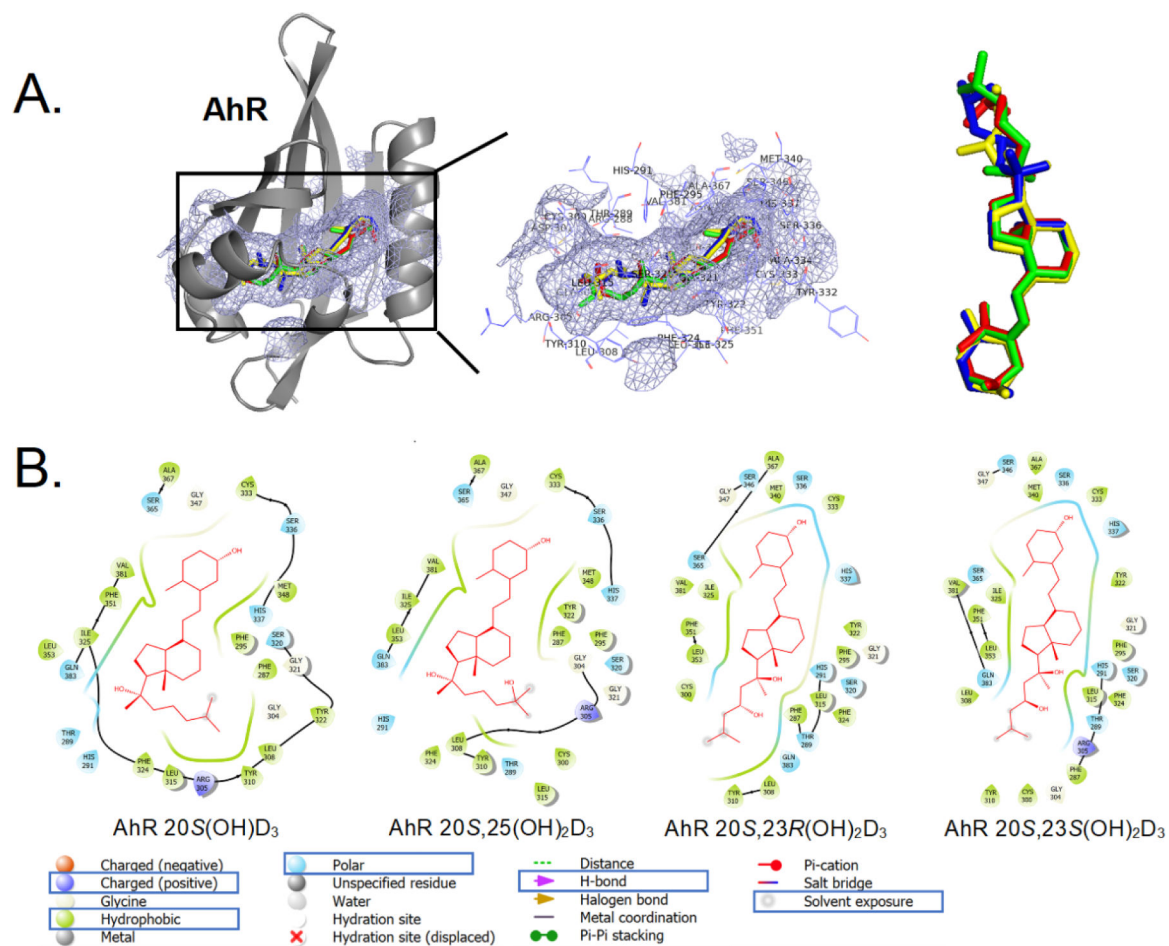


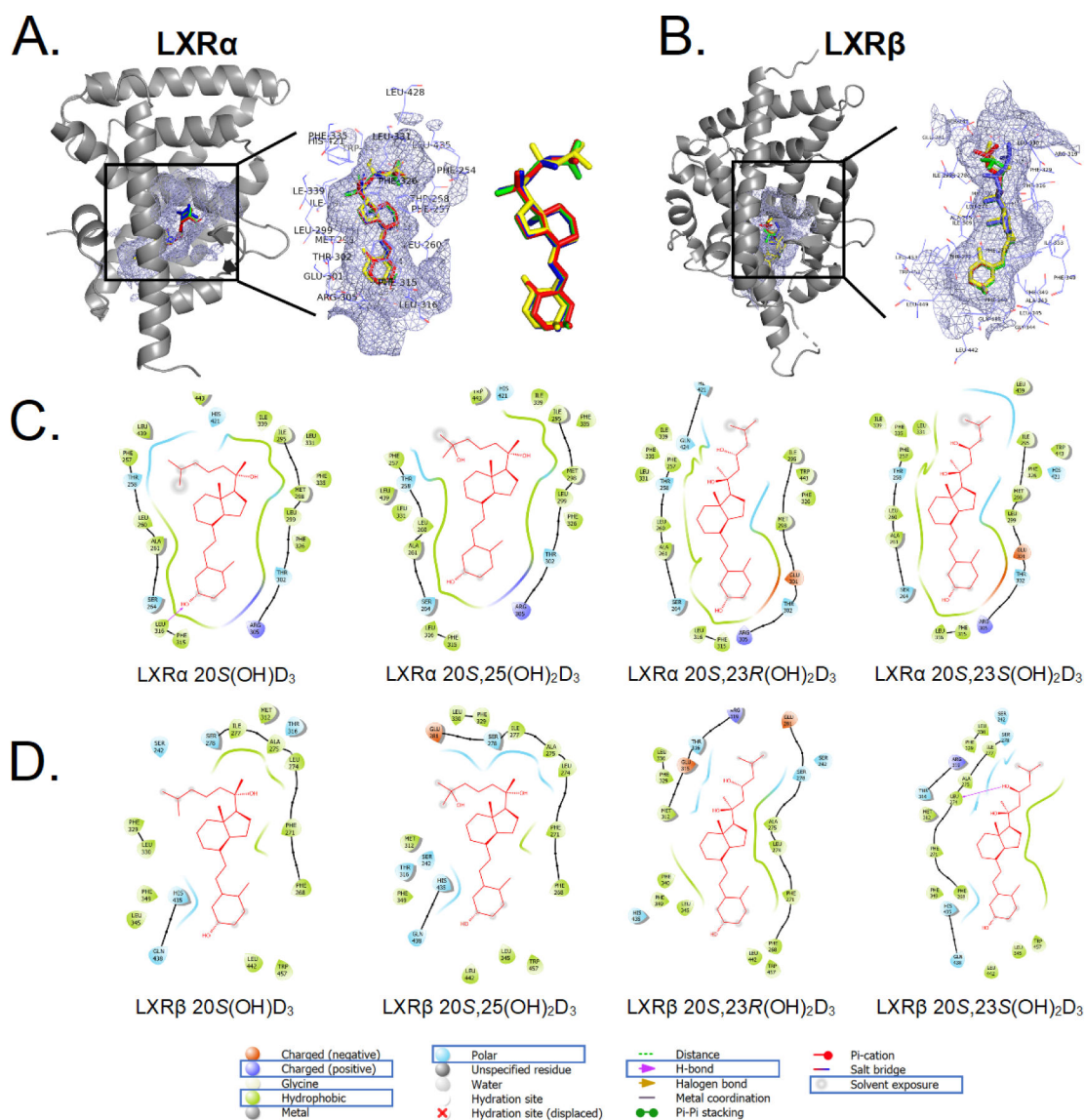
Figure 4. Stimulation of keratinocyte differentiation by 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ in comparison to 1 α ,25(OH)₂D₃. Human neonatal epidermal keratinocytes were treated for 8 (A), 24 h (B) or 48 h (C, D) with 0.1, 1, 10, or 100 nM secosteroid or 0.1% ethanol (negative control). A and B: relative gene expression studies with cyclophilin B used as internal control and presented as fold change vs vehicle control. C, D: protein expression studies. Values are presented as percentage of fluorescence positive cells (C) or as mean fluorescence per cell (% of all cells (D)). The values in D were calculated as mean green fluorescence vs DAPI blue fluorescence measured in ImageJ. E, F: Representative fluorescence (green) of cells stained with antibodies against involucrin (E) or catalase (F). Nuclei (blue) are stained with DAPI. Data are means \pm SD (n = 3) and were analyzed using Student's *t*-test with $p < 0.05$ (*), $p < 0.01$ (**) or $p < 0.001$ (***)

**Figure 5.**

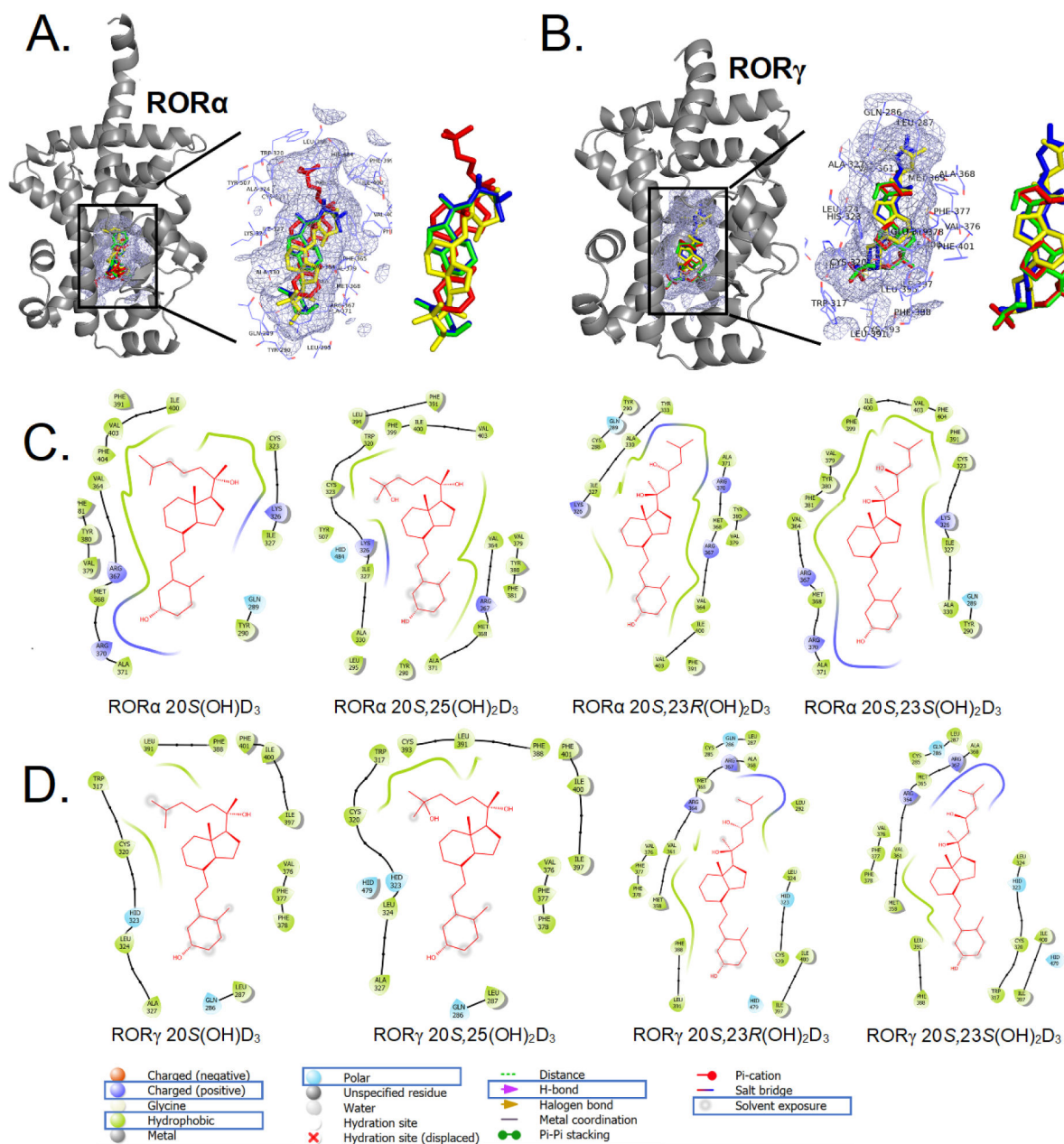
Interaction of 20S(OH)₂D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with nuclear receptors. A: ligand induced translocation of the VDR-GFP from the cytoplasm to nucleus. B: stimulation of AhR binding activity by secosteroids using AhR Reporter Cells including the luciferase reporter gene linked to an AhR-responsive promoter (Indigo Biosciences, State College, PA). C: ligand induced stimulation of CYP24A1 gene expression after 8 h of incubation. D: ligand stimulated expression of CYP1A1 and CYP1B1 genes. E: LXRα. F: LXRβ binding assay using LanthaScreen TR-FRET coactivator kit (Thermo Fisher Scientific, Inc., Waltham, MA). Data represent means ± SE where **p* < 0.05, ***p* < 0.01, ****p* < 0.001 and *****p* < 0.0001 at student *t*-test; #*p* < 0.05, ##*p* < 0.01, ###*p* < 0.001 and ####*p* < 0.0001 at one way ANOVA. For A (n = 4), B (n = 3), C and D (n = 3), E and F (n = 4). Gene expression was measured after 8 h of keratinocytes treatment with secosteroids and presented as fold change vs ethanol (0.1%) control and with GADPH (C) and cyclophilin (D) serving as internal controls.

**Figure 6a.**

The binding pattern of selected vitamin D₃ derivatives with AhR. A: 3D binding modes for 20S(OH)D₃ (green), 20S,25(OH)₂D₃ (red), 20S,23S(OH)₂D₃ (blue) and 20S,23R(OH)₂D₃ (yellow) in ligand binding domain of AhR (cartoon in gray). The binding pocket is shown as a light blue meshing area. B: 2D interaction map of 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with AhR (image generated with Maestro (v12.4)).

**Figure 6b.**

The binding pattern of selected vitamin D₃ derivatives with LXRs. A: 3D binding modes for 20S(OH)D₃ (green), 20S,25(OH)₂D₃ (red), 20S,23S(OH)₂D₃ (blue) and 20S,23R(OH)₂D₃ (yellow) in ligand binding domain of LXR α (cartoon in gray). B: 3D binding modes for 20S(OH)D₃ (green), 20S,25(OH)₂D₃ (red), 20S,23S(OH)₂D₃ (blue) and 20S,23R(OH)₂D₃ (yellow) in ligand binding domain of LXR β (cartoon in gray) including a zoomed view. The binding pocket is shown as a light blue meshing area. C: 2D interaction map of 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with LXR α . D: 2D interaction map of 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with LXR β (image generated with Maestro (v12.4)).

**Figure 6c.**

The binding pattern of selected vitamin D₃ derivatives with RORs. A: 3D binding modes for 20S(OH)D₃ (green), 20S,25(OH)₂D₃ (red), 20S,23S(OH)₂D₃ (blue) and 20S,23R(OH)₂D₃ (yellow) in ligand binding domain of ROR α (cartoon in gray). B: 3D binding modes for 20S(OH)D₃ (green), 20S,25(OH)₂D₃ (red), 20S,23S(OH)₂D₃ (blue) and 20S,23R(OH)₂D₃ (yellow) in ligand binding domain of ROR γ (cartoon in gray) including a zoomed view. The binding pocket is shown as a light blue meshing area. C: 2D interaction map of 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with ROR α . D: 2D interaction map of 20S(OH)D₃, 20S,25(OH)₂D₃, 20S,23S(OH)₂D₃ and 20S,23R(OH)₂D₃ with ROR γ (image generated with Maestro (v12.4)).

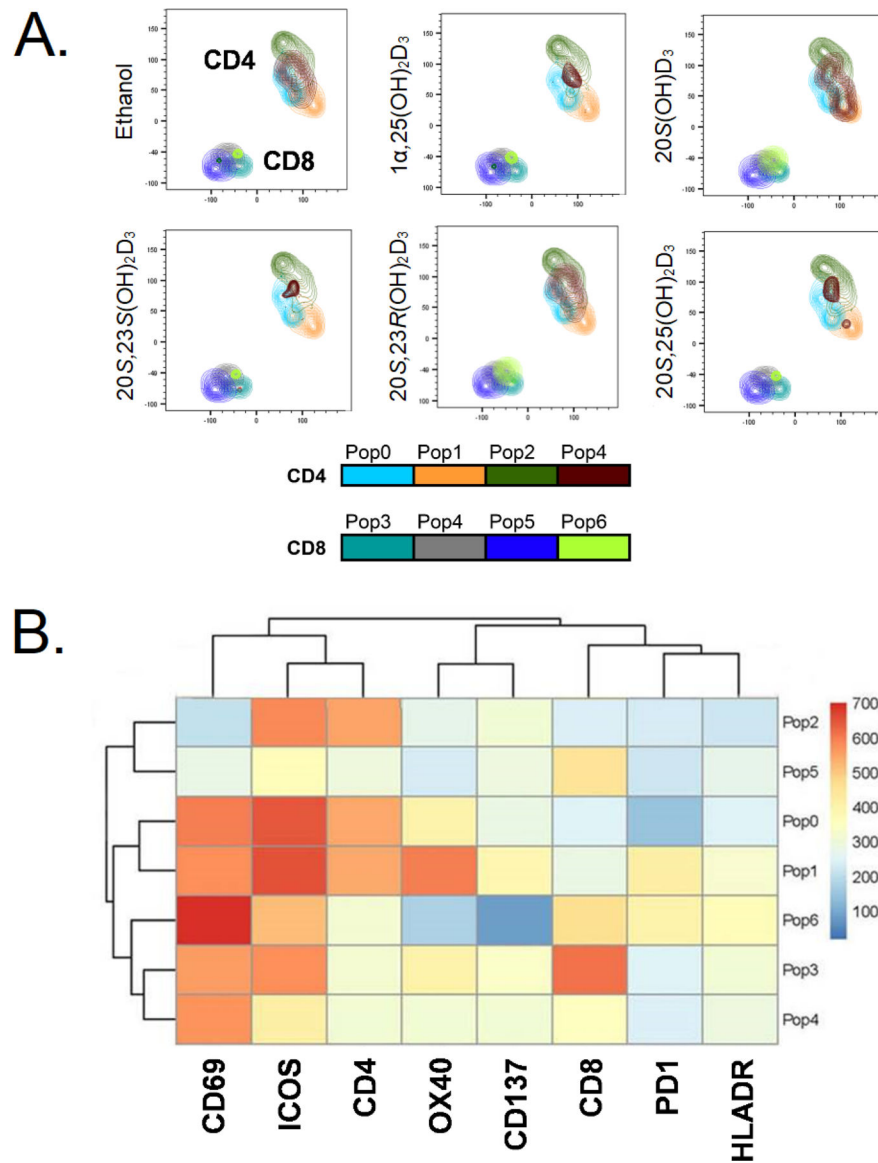
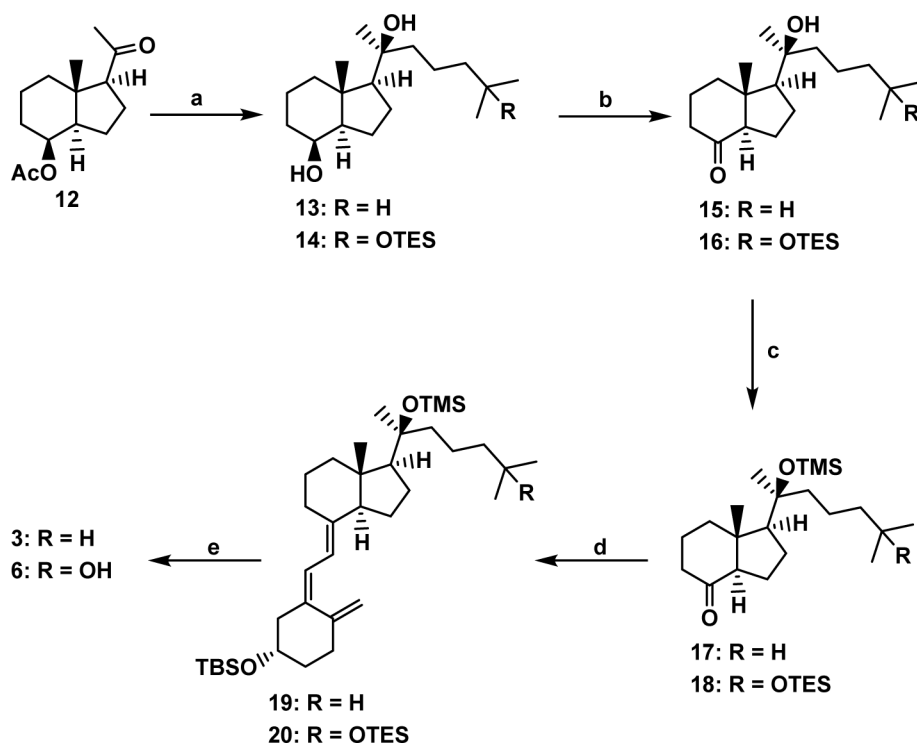
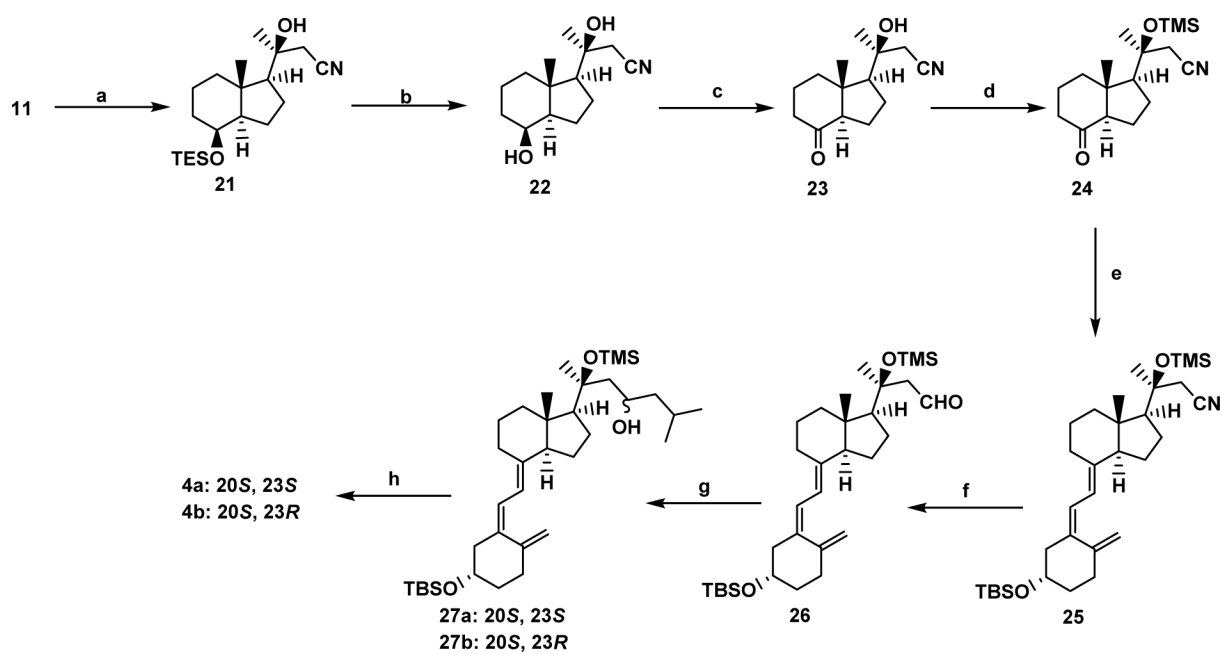


Figure 7. Differential and overlapping effects of secosteroids on activation of CD4 and CD8 T cells from human peripheral blood. A: FlowSOM analysis shows that the different vitamin D₃ secosteroids variably alter CD4 and CD8 T cell activation. B: Heat map of FlowSOM clusters representing the level of expression of activation induced markers (AIM) on CD4 and CD8 T cell population shown in A. T cells in peripheral blood from two healthy individuals were activated with anti-CD3 and anti-CD28 in the absence (ethanol) or presence of 10⁻⁷ M of each of the vitamin D₃ secosteroids for 24 h. The cells were stained for lineage markers (CD4, CD8, CD19, CD14) and a panel of activation induced markers (CD69, ICOS, OX40, CD137, PD1, HLADR) and expression data were collected using a flow cytometer. The gating strategy was live, single cell followed by CD4 and CD8 T cell populations. The data file output following following CD4 and CD8 T cell gating from each individual were concatenated into one file and subjected first to the dimensionality reduction algorithm

TriMAP, which clusters cell populations based on relatedness using three points. The data shows that CD4 and CD8 T cell populations cluster independently as expected. Following TriMAP analysis, we performed clustering with Self-Organizing Maps (FlowSOM) and the identified clusters were overlaid on to TriMAP clusters. Pop0 to Pop6 are subpopulations of cells based on expression levels of all markers shown in the heat map (Fig. 6B). Unbiased analyses using TriMap dimensionality reduction clustering followed by FlowSOM “R” algorithm-based clustering defined these subpopulations based on the expression levels of the markers. The algorithm-based clustering analysis takes into consideration the expression levels CD4, CD8 and all AIMS. The expression levels for each marker in specific FlowSOM populations is output as a heat map B. The heat map reflects the levels of expression of each marker within each subpopulation (Pop0 – Pop6) and is not by itself reflective of response to a vitamin D3 analogue.

**Scheme 1.**

(a) for **13**: Mg, Br(CH₂)₃CH(CH₃)₂, THF, 0 °C → rt, 85%; for **14**: Mg, Br(CH₂)₃C(CH₃)₂OTES, THF, 0 °C → rt, 80%; (b) TPAP, NMO, 4 Å MS, CH₂Cl₂, 0 °C → rt, **15**: 77%, **16**: 80%; (c) TMSOTf, 2,6-lutidine, CH₂Cl₂, -78 °C, **17**: 84%, **18**: 82%; (d) **9**, *n*-BuLi, THF, -78 °C, **19**: 81%, **20**: 91%; (e) TBAF, THF, rt, **3**: 84%, **6**: 75%.

**Scheme 2.**

(a) CH_3CN , LDA, THF, -78°C , 92%; (b) 4M HCl in dioxane, CH_2Cl_2 , -78°C , 98%; (c) TPAP, NMO, 4 Å MS, CH_2Cl_2 , $0^\circ\text{C} \rightarrow \text{rt}$, 95%; (d) TMSOTf, pyridine, 0°C , 74%; (e) **9**, *n*-BuLi, THF, -78°C , 87%; (f) DIBAL-H, CH_2Cl_2 , -78°C , 62%; (g) *i*-BuMgCl, THF, $0^\circ\text{C} \rightarrow \text{rt}$, 70%; (h) TBAF, THF, rt, 74%.

Table 1.Metabolism of chemically synthesized secosteroids by CYP27B1.^a

Secosteroid	1 α -hydroxylase activity (mol product/min/mol CYP27B1)
20,S(OH)D ₃	0.135 ± 0.005
20,S,23,R(OH) ₂ D ₃	0.117 ± 0.027
20,S,23,S(OH) ₂ D ₃	0.258 ± 0.020
20,S,25(OH) ₂ D ₃	7.98 ± 0.24

^aSecosteroids were incorporated into phospholipid vesicles at a ratio of 0.025 mol secosteroid/mol phospholipid and incubated with mouse CYP27B1 (0.2 μ M) for 20 min for all substrates except 20,S,25(OH)₂D₃ which was incubated for 5 min. Extracted samples were analyzed by reverse-phase HPLC. Data are mean \pm SD, n=3.

Table 2.

Molecular docking results for the complexes of vitamin D₃ hydroxy derivatives with selected nuclear receptors (kcal/mol).

Secosteroid \ Nuclear Receptor	VDR	ROR α	ROR γ	LXR α	LXR β	AhR	
Control	20(OH) Cholesterol	N/A	-11.4	-10.7	-10.0	-9.2	N/A
	25(OH) Cholesterol	N/A	-11.5	-10.9	-9.8	-9.8	N/A
	Indole acetic acid	N/A	N/A	N/A	N/A	N/A	-8.0
	1 α ,25(OH) ₂ D ₃	-11.8	-11.3	-10.7	-10.9	-11.0	-12.0
20S(OH)D ₃	-11.6	-10.7	-10.2	-10.2	-11.0	-11.9	
20S,25(OH) ₂ D ₃	-11.9	-10.8	-10.1	-10.2	-11.0	-11.6	
20S,23S(OH) ₂ D ₃	-11.4	-10.7	-10.1	-10.0	-11.1	-11.9	
20S,23R(OH) ₂ D ₃	-11.6	-10.7	-10.3	-9.9	-10.5	-11.8	