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Isoform selective PLD inhibition by novel, chiral 2,8-diazaspiro[4.5]decan-1-one derivatives

Alex G. Waterson^{a,e}, Sarah A. Scott^a, Nathan R. Kett^c, Anna L. Blobaum^{a,b}, H. Alex Brown^{a,c,d,e,f}, Craig W. Lindsley^{a,b,c,d,e,*}

^a Department of Pharmacology, Vanderbilt University Medical Center, Nashville, TN 37232, USA

^b Vanderbilt Center for Neuroscience Drug Discovery, Vanderbilt University Medical Center, Nashville, TN 37232, USA

^c Department of Biochemistry, Vanderbilt University, Nashville, TN 37232, USA

^d Department of Chemistry, Vanderbilt University, Nashville, TN 37232, USA

e Vanderbilt Institute of Chemical Biology, Vanderbilt University/Vanderbilt University Medical Center, Nashville, TN 37232, USA

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ABSTRACT

This letter describes the on-going SAR efforts to develop PLD1, PLD2 and dual PLD1/2 inhibitors with improved physiochemical and disposition properties as well as securing intellectual property position. Previous PLD inhibitors, based on a triazaspiro[4.5]decanone core proved to be highly selective PLD2 inhibitors, but with low plasma free fraction (rat, human $f_u < 0.03$), high predicted hepatic clearance (rat $CL_{hep} > 65 \text{ mL/min/kg}$) and very short half-lives *in vivo* ($t_{1/2} < 0.15$ h). Removal of a nitrogen atom from this core generated a 2,8-diazaspiro[4.5]decanone core, harboring a new chiral center, as well as increased sp³ character. This new core demonstrated enantioselective inhibition of the individual PLD isoforms, enhanced free fraction (rat, human $f_u < 0.13$), engendered moderate predicted hepatic clearance (rat $CL_{hep} ~ 43 \text{ mL/min/kg}$), improved half-lives *in vivo* ($t_{1/2} > 3$ h), and led to the first issued US patent claiming composition of matter for small molecule PLD inhibitors.

Phospholipase D (PLD) exists as two isoforms in mammals, coined PLD1 and PLD2, and is considered to be one of the major sources of the signal-activated second messenger phosphatidic acid (PtdOH). Recently, the development of isoform-selective PLD inhibitors, in combination with biochemical and molecular genetics studies, have suggested that PLD enzymes in mammalian cells and pathogenic organisms represent therapeutic targets for several human diseases.^{1–5} Work from multiple labs have demonstrated that PLD dysfunction and/or overexpression can be modulated by small molecule, allosteric inhibitors 1-6 (Fig. 1) with efficacy in preclinical models relevant to oncology, viral infections and CNS disorders.^{1–17} Despite the progress achieved in the past decade, small molecule tools with DMPK profiles suitable for robust in vivo proof of concept studies have remained elusive, and there have been no PLD inhibitor composition of matter patents (issued or published applications), further hindering translational potential for the target.¹⁻²² In this Letter, we describe the discovery and development of a new series of potent and selective PLD2 and dual PLD1/2 inhibitors from a series of 2,8-diazaspiro[4.5]decanones, that display enantioselective PLD isoform inhibition, provide improved DMPK profiles and enabled a robust intellectual property (IP) position. 23

For several years, the triazaspiro[4.5]decanone core of PLD inhibitors 3-6 was the major focus of optimization efforts, as both PLD2selective and dual PLD1/2 inhibitors could be realized within this core based on chiral α -methyl moieties along the ethyl linker chain (e.g., 5).^{7–14} Despite providing both peripherally restricted and highly CNS penetrant (brain-to-plasma Kps 0.75-1.48) PLD inhibitors, this series uniformly afforded compounds displaying high plasma protein binding (rat and human $f_{\rm u}$ < 0.03), high predicted hepatic clearance (rat $CL_{hep} > 65 \, mL/min/kg$) and very short half-lives in vivo ($t_{1/}$ $_2 < 0.15$ h).⁷⁻¹⁴ Work within a well-known SERM chemotype afforded only weak PLD inhibitors, but they proved to be universal (mammalian and bacterial).¹⁵ Thus, the ideal in vivo proof of concept tools were lacking, and the program also needed novel chemical matter to advance the target. Our strategy to potentially accomplish both goals focused on deletion of the N-aryl nitrogen atom in 7, and replacement with a stereogenic carbon center, as in 8 (Fig. 2). We were also interested to see the impact of a chiral α -methyl moiety on the linker with regards to

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^{*} Corresponding author at: Department of Pharmacology, Vanderbilt University Medical Center, Nashville, TN 37232, USA.

E-mail address: craig.lindsley@vanderbilt.edu (C.W. Lindsley).

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Ŵе Ô R VU0359595 (2) VU0364739 (3) halopemide (1) PLD1 IC₅₀ = 3.7 nM PLD1 IC₅₀ = 1,500 nM PLD1 IC₅₀ = 21 nM PLD2 IC₅₀ = PLD2 IC₅₀ = 6,400 nM 20 nM PLD2 IC₅₀ = 300 nM 1,700-fold PLD1 selective 75-fold PLD2 selective dual inhibitor

Fig. 1. Halopemide (1) was the progenitor of all PLD inhibitors, and a dual PLD1/2 inhibitor. Optimization efforts led to isoform-selective PLD inhibitors: 2 (1700-fold PLD1 selective), 3 (75-fold PLD2 selective), 4 (53-fold PLD2 selective), 5 (dual PLD1/2 inhibitor) and 6 (> 80-fold PLD2 selective).

ML299 (5)

PLD1 $IC_{50} = 6 nM$

PLD2 IC₅₀ = 20 nM

dual inhibitor

Mé



Fig. 2. Envisioned optimization strategy for 7 to improve disposition and structural novelty with analogs 8.

PLD isoform inhibition (e.g., the 'magic methyl' effect)²⁴ in combination with the phenyl bearing stereocenter, and if previous SAR for **3–6** would be maintained.^{7–14}

ML298 (4)

PLD1 IC₅₀ = 20,000 nM

53-fold PLD2 selective

355 nM

PLD2 $IC_{50} =$

Fortunately, Thomas and co-workers form Roche had previously reported on a racemic synthesis of a 4-fluorophenyl 2,8-diazaspiro[4.5] decanone core,²⁵ and had provided details for the assignment of the individual enantiomers. Thus, we elected to follow their route and first evaluate both enantiomers of this core for PLD inhibition. As prescribed (Scheme 1), deprotonation of **9** with LDA followed by trapping with *trans*-1-fluoro-4-(2-nitrovinyl)benzene afforded the desired Michael adduct **10** in 90% yield. Reduction of the nitro moiety with Ra-Ni afforded primary amine **11** in 86% yield. Finally, closure to the racemic spiro- γ -lactam (\pm)-**12** was facilitated by refluxing **11** in toluene overnight; however, some deprotection of the Boc was noted, so the mixture was reprotected to afford Boc-protected (\pm)-**12** in 83% yield. Preparative chiral separation (2–3 g batches) was performed on a Gilson 215 Preparative HPLC with pumps capable of 200 mL/min using a Chiraltech OJ 20u, 50 × 250 mm column at a flow rate of 177 mL/min (95% hexanes and 5% isopropyl alcohol) to afford (*S*)-12 and (*R*)-12, and in agreement with the work of Thomas.²⁵

Once in hand, (*S*)-12 and (*R*)-12 were rapidly elaborated into putative PLD inhibitors (*S*)-14 and (*R*)-14 in a two-step sequence (Scheme 2).^{7–14} Reductive amination with *tert*-butyl 2-oxoethylcarbamate and (*S*)-12 and (*R*)-12 provided (*S*)-13 and (*R*)-13, respectively in 79–85% yield after Boc deprotection. Acylation of the primary amine with the requisite acid chloride delivered (*S*)-14 and (*R*)-14 in yields ranging from 58 to 95%. To assess the impact of the (*S*)-methyl group on the side chain, a similar two step route was employed (Scheme 3) wherein (*S*)-*tert*-butyl 1-oxo-2-propanylcarbamate is substituted for *tert*-butyl 2-oxoethylcarbamate with comparable yields at each step.

As shown in Table 1, clear enantioselective inhibition of the PLD isoforms was noted, with all (*R*)-14 analogs typically > 10-fold weaker at inhibiting both PLD1 and PLD2 versus the (*S*)-14 analogs. For example, (*S*)-14a was a 70 nM PLD 2 inhibitor with ~17-fold selectivity versus PLD1, while (*R*)-14 was > 10-fold weaker at PLD2 ($IC_{50} = 810$ nM). This SAR trend was conserved across all enantiomeric



ML395 (6)

PLD1 IC₅₀ >30,000 nM

PLD2 IC₅₀ = 360 nM

>80-fold PLD2 selective



Scheme 1. Synthesis of 4-fluorophenyl 2,8-diazaspiro[4.5]decanone cores **12**.^a ^aReagents and conditions. (a) *trans*-1-fluoro-4-(2-nitrovinyl)benzene, LDA, THF, -40 °C, 2 h, 90%; (b) H₂, Ra-Ni, EtOH, 50 °C, 20 h, 86%; (c) i) toluene, reflux; ii) Boc₂O, DIPEA, dioxane-H₂O, 0 °C, 2 h, 83%; d) i) Gilson 215 Preparative HPLC with pumps capable of 200 mL/min using a Chiraltech OJ 20u, 50 × 250 mm column at a flow rate of 177 mL/min (95% hexanes and 5% isopropyl alcohol. (*S*)-**12** is first eluting peak; ii) TFA, CH₂Cl₂, rt, 5 h, 95%.



Scheme 2. Synthesis of 2,8-diazaspiro[4.5]decanone-based PLD inhibitors **14**.^a Reagents and conditions: (a) i) *tert*-butyl 2-oxoethylcarbamate, MPB(OAC)₃H, DCM, rt; ii) 4.0 M HCl/dioxane, DCM, MeOH, rt, 79–85%; (b) RCOCl, DIEA, DMF, rt, 58–95%.



Scheme 3. Synthesis of 2,8-diazaspiro[4.5]decanone-based PLD inhibitors 16.^a ^aReagents and conditions: (a) i) (*S*)-*tert*-butyl 1-oxo-2-propanylcarbamate, MPB (OAc)₃H, DCM, rt; ii) 4.0 M HCl/dioxane, DCM, MeOH, rt, 70–82%; (b) RCOCl, DIEA, DMF, rt, 55–92%.

pairs of (*S*)- and (*R*)-14 derivatives. Thus, further optimization will maintain the (*S*)-configuration at the aryl bearing stereogenic center. The SAR trend within 3–6 by which a chiral (*S*)-methyl moiety on the linker not only improved PLD activity, but also enhanced PLD1 inhibitory activity was retained in the diazaspiro[4.5]decanone series.^{7–14} For example, addition of the (*S*)-methyl moiety as in (*S*)-16a increased PLD1 activity ~9-fold over (*S*)-14a, providing an ~10-fold PLD1-prefering inhibitor. With the analogous (*R*)-16a, activity was increased at PLD1, but with a significant loss of inhibitory activity at PLD2. This enantiospecific modulation of PLD activity was also conserved within the entire series, with the (*S*)-methyl group engendering enhanced PLD1 inhibition with a concomitant loss of PLD2 activity. Notable examples include (*S*)-16b, a potent dual PLD1 and PLD2 inhibitor (IC₅₀s of 1 nM and 12 nM, respectively) and (*S*)-14b, a 25-fold PLD2 preferring inhibitor, which converts to a dual PLD1/PLD2 inhibitor (*S*)-16d







Compound	R	PLD1 IC ₅₀ (nM) ^a	PLD2 IC ₅₀ (nM) ^a
(S)-14a		1200	70
(R)- 14a	i i i i i i i i i i i i i i i i i i i	6400	810
(S)- 16a		16	150
(R)- 16a		320	4000
(S)- 14b		21	2.5
(R)-14b		320	43
(S)-16b		1	12
(R)- 16b	H H	16	140
(S)-14c		1350	60
(R)- 14c		12,000	450
(S)-16c	Y V V	25	115
(R)- 16c		200	700
(S)-14d	¥ ~ ~	5500	220
(R)- 14d	F	> 20,000	3175
(S)-16d	F	250	210
(R)- 16d	F	1700	2800
(S)- 16e		820	26

^bPLD2 cellular assay in HEK293-gfp-PLD2 cells. IC₅₀ values are the average of three independent measures (n = 3).

^a Cellular PLD1 assay in Calu-1 cells.

upon incorporation of the (*S*)-methyl group. These data proved very exciting, confirming that a single chemotype can provide PLD1-preferring, PLD2-preferring and dual PLD1/PLD2 inhibitors (Fig. 3) based solely on the inclusion of either one or two easily established and constructed chiral centers. These data also supported the first, and only to date, issued U.S. composition of matter patent for PLD inhibitors.²³

We next evaluated if these novel PLD inhibitors offer advantages in terms of disposition relative to the earlier generation **2–6**, which are plagued by high plasma protein binding (rat and human $f_u < 0.03$), high predicted hepatic clearance (rat CL_{hep} > 65 mL/min/kg) and very short half-lives *in vivo* ($t_{1/2} < 0.15$ h). As shown in Table 2, analogs **14** and **16** uniformly showed improved free fraction (in both rat and human plasma, $f_us > 0.10$) and (*S*)-**14b** and (*S*)-**14d** displayed improved predicted hepatic clearance in rat (41.2 and 47.2 mL/min/kg, respectively) and human (10.7 and 10.1 mL/min/kg, respectively).



Fig. 3. Representative PLD1 (Calu-1) and PLD2 (HEK293-PLD2) concentration-response-curves for (S)-14b, a PLD2-preferring inhibitor and (S)-16d, a dual PLD1/PLD2 inhibitor.

Table 2

In vitro DMPK data for analogs (S)-14 and (S)-16.

Property	(S)-14a	(S)-14b	(S)-14c	(S)-14d	(<i>S</i>)-16b	(<i>S</i>)-16e
MW (dal) cLogP	419.5 2.87	434.2 2.29	446.2 2.68	413.4 2.84	448.5 2.73	459.5 3.31
TPSA (Å ²)	61.4	73.4	73.8	92.9	73.4	61.4
In vitro PK parameters Rat CL _{HEP} (mL/min/	60.7	41.2	63.8	47.2	53.4	61.6
kg) Human CLump (mL/	61 7	10.7	17 2	10.1	14.0	174
min/kg)	01.7	10.7	17.2	10.1	14.0	17.4
Rat f _u plasma	0.160	0.198	0.314	0.246	0.130	0.061
Hum $f_{\rm u}$ plasma	0.138	0.124	0.122	0.108	0.136	0.045
Rat $f_{\rm u}$ brain	0.013	0.004	0.026	0.027	0.004	0.003

Based on the improved predicted hepatic clearance of (S)-14b and (S)-14d, we evaluated both in a rat IV PK cassette study (0.25 mg/kg per compound in EtOH/PEG400/DMSO vehicle). Both compounds displayed suprahepatic clearance in vivo (CLps of 340 and 250 mL/min/kg for (S)-14b and (S)-14d, respectively), and thus a lack of an in vitro:in vivo correlation (IVIVC). However, both compounds showed improved half-lives (*t*_{1/2}s of 3.63 and 3.2 h, for (*S*)-14b and (*S*)-14d, respectively) driven by large volumes of distribution at steady state (Vss of 80 and 56 L/kg for (S)-14b and (S)-14d, respectively). Thus, despite attractive cLogPs, these new inhibitors were clearly deporting in some tissue and being reabsorbed into plasma, providing a sustained half-life and mean residence time. Based on these data and associated concerns regarding accumulation with chronic dosing and the subsequent potential for tox findings, this series could not be further progressed. Still, this new series of 2,8-diazaspiro[4.5]decanones provided improvements in multiple domains over historical PLD inhibitors 2-6.

In summary, we employed a scaffold-hopping strategy, moving from the classic triazaspiro[4.5]decanone-based PLD inhibitors to a novel 2,8-diazaspiro[4.5]decanone core, harboring a stereogenic center. This new core demonstrated enantioselective inhibition of the individual PLD isoforms (providing access to PLD1, PLD2 and dual PLD1/PLD2 inhibitors in the same scaffold, along with enhanced plasma free fraction (rat, human $f_u < 0.13$), engendered moderate predicted hepatic clearance (rat $CL_{hep} \sim 43 \text{ mL/min/kg}$), and improved half-lives *in vivo* ($t_{1/2} > 3 \text{ h}$). Further, this novel chemical matter led to the first issued US patent claiming composition of matter for small molecule PLD inhibitors. While this new series displayed robust and tractable SAR, the extremely high volume of distribution *in vivo* precluded further advancement. Follow-up efforts continue, and work is in progress to develop optimal *in vivo* tool compounds that selectively inhibit either PLD1 or PLD2 as well as clinical candidates.

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