

Development of DGK Inhibitors, hit validation of initial HTS

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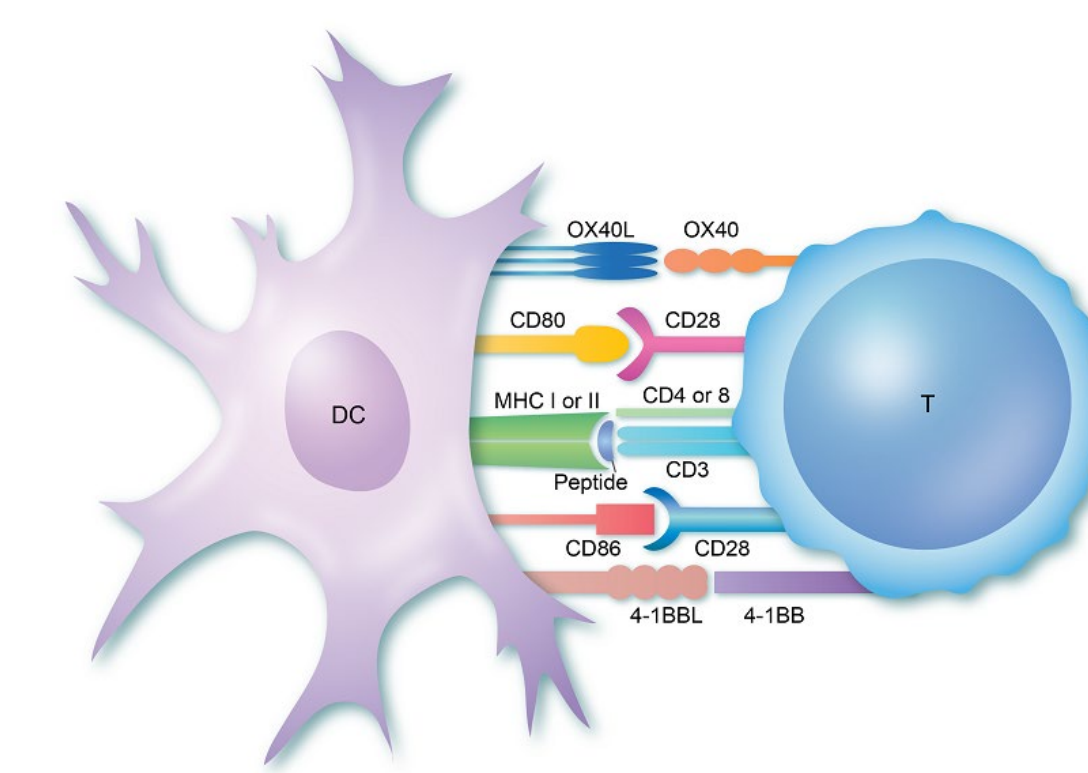
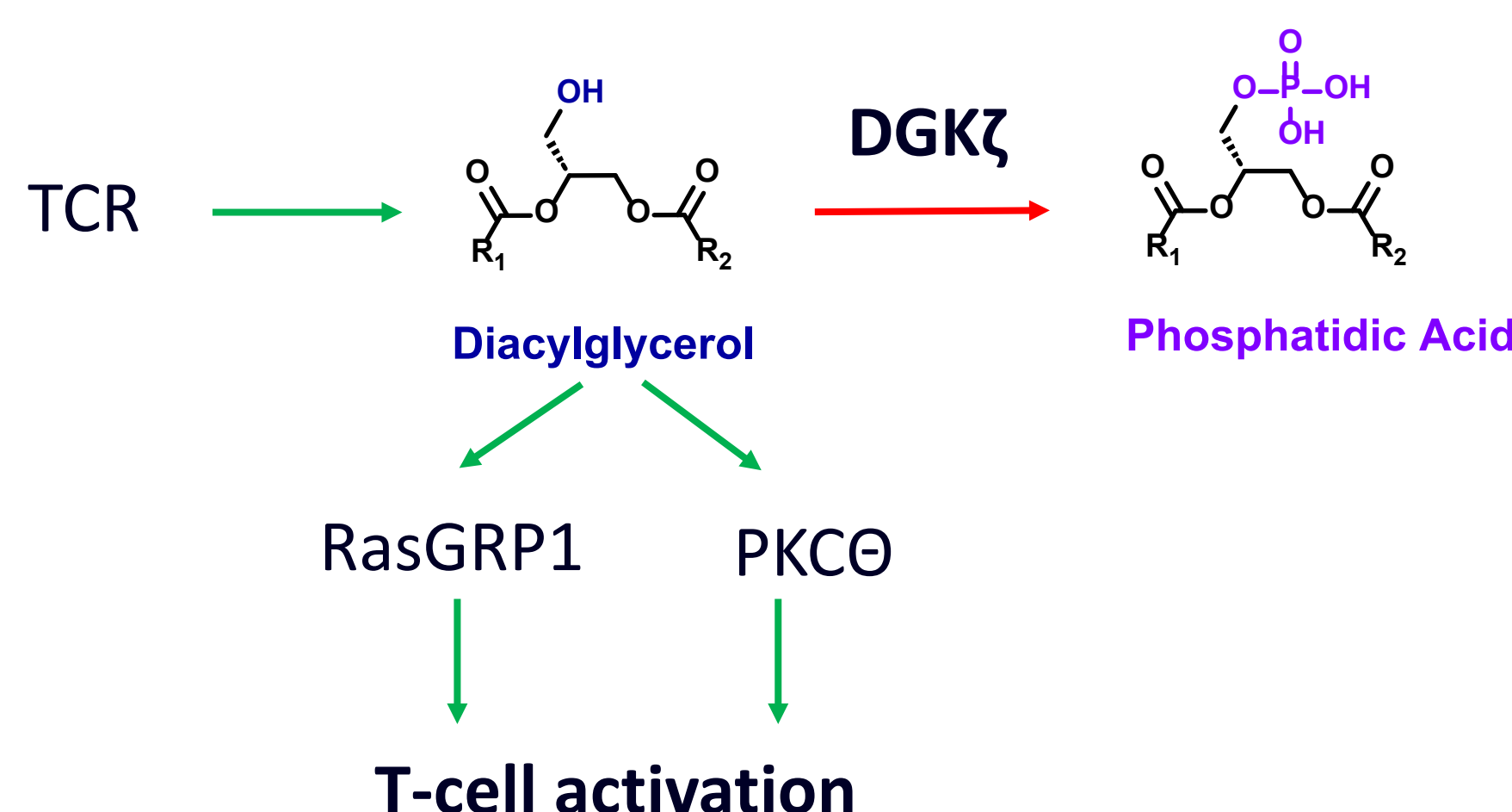
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TARGET HYPOTHESIS

Disease setting: Immune oncology, solid tumours.

- **DAG regulates immune cell signalling and activation** (T-cells and NK cells).
- Diacylglycerol kinase zeta (**DGK ζ**) catalyses the phosphorylation of DAG to PA.
- DGK has 10 isoforms, ζ and α are those mostly expressed in the immune system.
- **Limiting the pool of DAG effects a positive signalling through T-cells.**



Schematic of early T cell activation through binding to a dendritic cell (DC). Subsequently, co-stimulation occurs through DC-bound CD86, CD80, OX40L and 4-1BBL. This induces full activation and effector function in the T cell.¹

OBJECTIVES and METHODOLOGY

- Tumour evasion of immune responses linked to **DGK ζ** and **DGK α** deregulation.
- **Inhibition** of both DGK ζ and DGK α to maintain higher levels of DAG, promote T cell proliferation and anti-tumour activity.
- Thus, **activating the immune response in "cold" tumours, resulting in activation of cytokine and inflammatory responses to stimulate the immune system to kill tumours.**
- Potent mixed inhibitors targeting DGK ζ and DGK α identified at the onset of the project, the specific effects of selective inhibition remained unexplored.

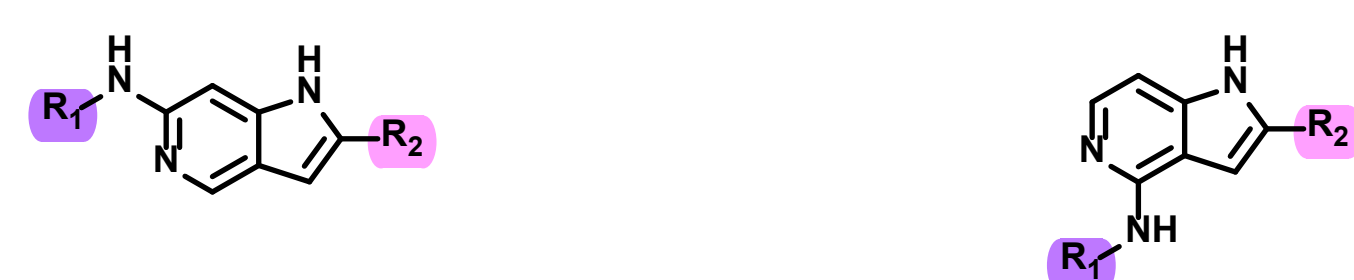
- Finding a "first-in-class" oral inhibitor of DGK ζ .
- Screening of >100k compounds by HTS (done at Charles River).



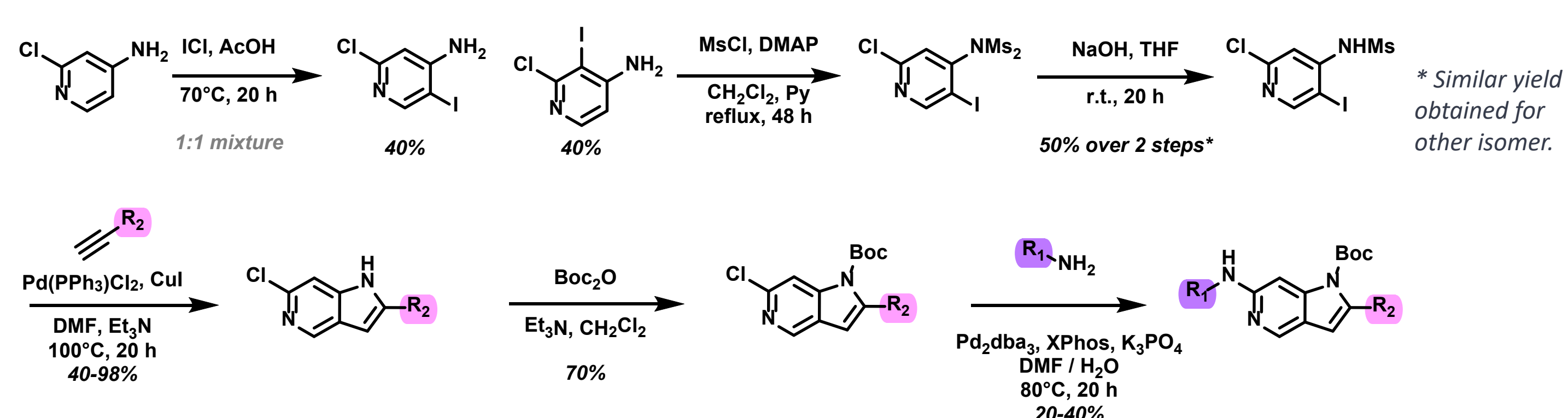
- 2 Series expanded using in-house developed **ADP-glo assay** and *in vitro* cellular testing.

AZAINDOLE SERIES

- From initial triage, cluster of 40 compounds with **IC₅₀ < 10 μ M**.
- SAR-study around two regioisomers was carried out.



Library Synthesis:



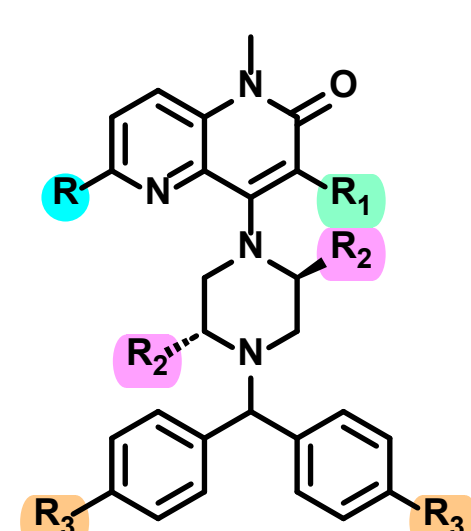
SAR:

R ₁	R ₂	IC ₅₀ (μ M)
Me	H	49
OMe	H	19
Cl	H	20
CF ₃	H	204
Me	Me	37
Me	OMe	37
Me	Me	32
Me	OMe	34
OMe	H	21
OMe	Me	18*
OMe	OMe	23
OMe	Me	37
OMe	OMe	32
OMe	OMe	34
OMe	OMe	148
OMe	OMe	165
OMe	OMe	20*
OMe	OMe	95
OMe	OMe	>100
OMe	OMe	58
OMe	OMe	61
OMe	OMe	84
OMe	OMe	9
OMe	OMe	44
OMe	OMe	85
OMe	OMe	42
OMe	OMe	70
OMe	OMe	76
OMe	OMe	71
OMe	OMe	30
OMe	OMe	9
OMe	OMe	6
OMe	OMe	23
OMe	OMe	23
OMe	OMe	21

* Original hits from HTS, values are for retest in ADP-Glo assay (in house).

COMPETITORS

- BMS has 2 optimized leads. Very potent but not selective. IC₅₀ measured using biochemical lipid kinase assay.²



BMS-502 - R = CN, R₁ = NO₂, R₂ = H, R₃ = F
IC₅₀ DGK ζ = 0.002 μ M, α = 0.005 μ M

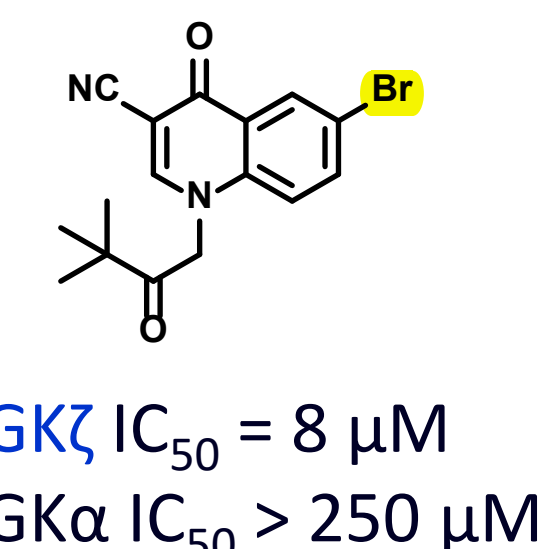
BMS-332 - R = CN, R₁ = H, R₂ = Me, R₃ = Me
IC₅₀ DGK ζ = 0.008 μ M, α = 0.009 μ M

CONCLUSION

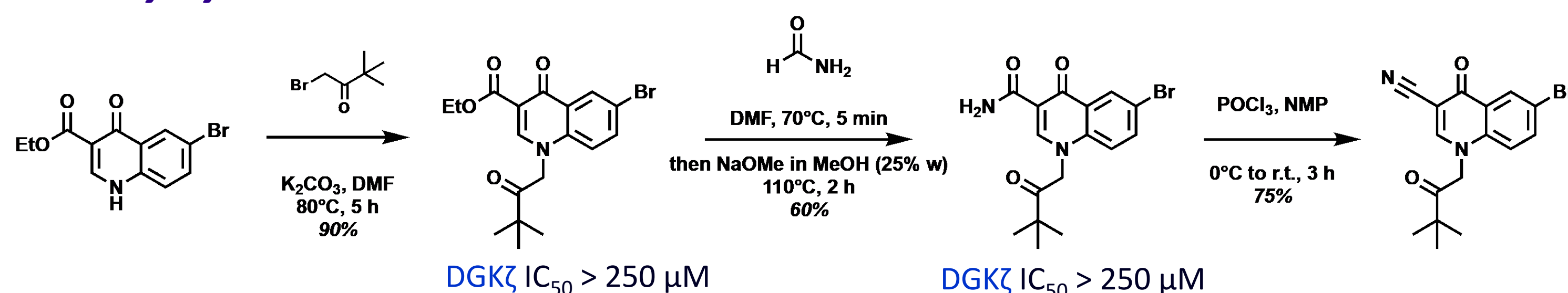
- Extensive synthetic efforts to expand both series with SAR studies.
- **Azaindole series** hindered by lengthy synthesis and poor reactivity (e.g. Buchwald-Pd coupling in last step).
- **Dihydroquinolinone series** widely expanded and explored, **12-fold potency gain with ethynyl-linker** (Potential *hit-to-lead*).
- Unfortunately, no DGK α IC₅₀ for all compounds due to assay format / protein production / purification.
- Competitors are in clinic with very potent (but not selective) candidates.

DIHYDROQUINOLINE SERIES

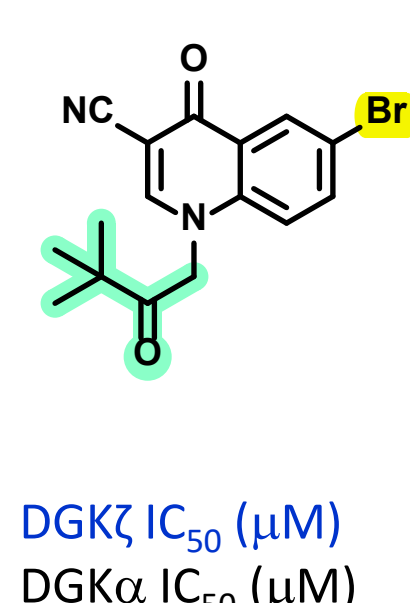
- Singleton choice with IC₅₀ < 10 μ M.
- **Br atom offers excellent synthetic vector for growing the fragment.**
- SAR around this region and around the "southern" part of the hit is shown below.



Library Synthesis:



SAR:



R ₁	R ₂	IC ₅₀ (μ M)
Me	H	39
OMe	H	4.5
OMe	Me	>250
OMe	OMe	0.61
OMe	OMe	46
OMe	OMe	0.41
OMe	OMe	4.2
OMe	OMe	14
OMe	OMe	>250

R₁ substitution highly tolerated in *para* position, especially E.D.G.

R ₁	R ₂	IC ₅₀ (μ M)
OMe	H	1
OMe	H	33
OMe	Me	1
OMe	Me	9
OMe	OMe	4.5
OMe	OMe	>250
OMe	OMe	4.5
OMe	OMe	>250

R ₃	IC ₅₀ (μ M)
Me	4
Et	4
iPr	36
Bn	>250
Ph	>250
Me	>250
Et	>250
iPr	>250
Bn	>8
Ph	99

- Other R₂ substituents not tolerated (i.e. pyridine, pyrimidines, bicycles, sat.)

R ₂	IC ₅₀ (μ M)
Me	67
OMe	n.d.
OMe	49
OMe	n.d.
OMe	25
OMe	157
OMe	>250
OMe	6

- Expansion of 5-memb set, tolerability and selectivity.

- SAR on "southern" part = large bulky ketone helps with potency.
- Better interactions to be found?

ACKNOWLEDGEMENTS

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