Perforin (PRF) is a calcium-dependent pore-forming glycoprotein found within granules of cytotoxic T lymphocytes (CTL) and natural killer (NK) cells. These cytotoxic effector cells play an important role in immune-surveillance and homeostasis via a granule exocytosis pathway responsible for the elimination of necrotic/virus-infected cells and intracellular pathogens. Numerous autoimmune diseases (e.g., insulin-dependent diabetes) and therapy-induced conditions (e.g., allograft transplantation rejection, graft-vs-host disease) adopt a synonymous biochemical pathway. An integral part of this biological cascade is perforin. 1,2 Evolution from a furan-containing high-throughput screen (HTS) hit (1) resulted in isobenzofuran-1(3H)-one (2) as a potent inhibitor of the function of both isolated perforin protein and perforin delivered in situ by intact KHYG-1 NK cells. In the current study, structure–activity relationship (SAR) development towards a novel series of diarylthiophene analogues has continued through the use of substituted-benzene and -pyridyl moieties as bioisosteres for 2-thioxoimidazolidin-4-one (A) and isobenzofuranone (C) scaffolds. The resulting compounds were tested for their ability to inhibit perforin lytic activity in vitro. Carboxamide (23) shows a 4-fold increase over (2) in lytic activity against isolated perforin and provides good rationale for continued development within this class.

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candidates for further in vivo studies showed some toxicity toward whole NK cells. It is therefore likely that toxicity would also be observed in the immunocompromised mice required for an efficacy study. Also desirable would be the identification of new compounds with significantly improved potency to deliver a lower efficacious dose and better control of potential dose-related toxicity. Finally, while the thioxoimidazolidinone series passed various pan-assay interference compound (PAINS) filters, they still contain a potentially reactive Michael acceptor and exist as an inseparable (and interconverting) mixture of E- and Z-isomers.

To address these aforementioned issues the need for a 2-thioxoimidazolidin-4-one bioisosteric replacement was apparent and herein we report on the latest developments towards an exciting new class of nontoxic inhibitors of perforin function that show improved inhibition of the function of both isolated perforin protein and perforin delivered in situ by intact KHYG-1 NK cells.

**Synthesis: Target compounds 6–23 (Scheme 1)** were obtained in a linear four-step sequence starting with the conversion of commercially available phenyl-boronates/boronic acids in the presence of catalytic PdCl₂(dppf), afforded the respective diaryliothiophene targets 6–23 in generally good yields. Isoindolinone-based versions of 23 (23a and 23b) were also prepared in exactly the same manner, using the isoindolin-1-one and 2-methylisoindolin-1-one-based analogues of key bromide 5 (5a and 5b; Scheme 1).

Analogous methodology was adopted to generate secondary and tertiary carboxamide targets 29 and 30, by first reacting iodide 24 with 2-bromothiophene in the presence of AgNO₃/KF (Scheme 2). The afforded bromide 25 was then converted to carboxamide intermediates 26 and 27 through activation of the carboxylic acid to a pentafluorophenyl (PFP) ester followed by subsequent addition of the corresponding amine (methylamine or dimethylamine) in pyridine for 2–3 h at room temperature. Key boronate 28 was prepared in multi-gram quantities from bromide 3 (Scheme 1) and subsequently used in a Suzuki coupling with intermediates 26 and 27 to furnish carboxamide targets 29 (75%) and 30 (57%), respectively.

Compounds 31 and 32, where the carboxamide side-chain was elaborated further by incorporating a series of basic side-chains, were accessed by applying the same chemistry as summarised in Scheme 1. A final step Suzuki coupling of key intermediate bromide 5 with the corresponding phenyl boronate (as according to the compounds of Table 1) gave 31 and 32 in good to modest yields, respectively, (61% and 42%).

With the chemistry now established for the mono-substituted A-ring targets shown in Schemes 1 and 2, a series of di-substituted targets were investigated. Phenyl iodides 33–35 were prepared via

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**Figure 1.** The development of new diaryliothiophene inhibitors of perforin.
esterification from their native carboxylic acids in the presence of EtOH and catalytic H₂SO₄ using conventional methodology (Scheme 3). The sequential step was carried out as for carboxylic acid analogue 25 (refer to Scheme 2) affording ethyl esters 36–38, these in turn were then coupled with key boronate ester 28 via a Suzuki reaction to give diaryliothiophene esters 39–41 in good to excellent yields. Hydrolysis of the resulting esters with 2 M HCl in MeOH under reflux conditions gave bis-carboxylic acid intermediates 42–44, respectively. Subsequent dehydration to isobenzofuran-1(3H)-one precursors 45–47 was achieved quantitatively using a mixture of TFA(CH₂Cl₂ 1:1) at room temperature for approximately 3–4 h. Target compounds 48–57 were then synthesised utilising a variety of amide-coupling strategies (Scheme 3). Carboxamide 48 was prepared successfully in 57% yield going via the analogous PFP-active ester intermediate as employed for carboxamides 26 and 27 (refer to Scheme 2). Compounds 49–54, which contain various extended neutral and basic side-chains appended to the 4-carboxamide moiety, were accessed through treatment of the corresponding acid precursor with DCC and catalytic HOBt at elevated temperatures. Target compounds 55 (48%), 56 (80%), and 57 (38%), all of which contain a 3-substituted-phenyl primary amine, were accessed through an EDCl/HOBt-mediated amide-coupling in the presence of anhydrous DMF at 0 °C to room temperature over 1–2 h. 

Compounds 58–63 (Scheme 1) encompass a variety of A-ring moieties (such as an alcohol, a methanethiol and various sulfonamides). These were all accessed through a key Suzuki coupling step from key intermediate 5 with the corresponding substituted boronate, as according to the compounds of Table 1. Isobenzofuran-1(3H)-one dimer (71) was prepared in the same manner.

Bromide 64 was commercially available and was converted to iodide 65 via a copper-mediated halogen exchange reaction as adapted from literature procedure 13 (Scheme 4). Boronate 66 was synthesised from corresponding bromide intermediate 5 in the presence of bis(pinacolato)diboron and KOAc and subsequently coupled with iodide 65 utilising Suzuki methodology to yield pyridyl carboxamide 67 in low yield.

Finally, the preparation of 4-pyridyl-containing target 70 began with the Suzuki-coupling of 4-bromopyridine 68 with thiophene-2-boronic acid, as described by Effenberger et al., 19 to furnish 4-(2-thienyl)pyridine 69 in high yield. Isobenzofuran-1(3H)-one 4 was then installed through treatment with AgNO₃ and KF in the presence of a palladium-complex, as previously discussed, to afford 4-pyridyl derivative 70 in 27% yield (Scheme 1).

Structure–activity relationships: Based on our previous studies described above, we hypothesised that selected substitutions on a benzene or pyridyl ring could potentially occupy the same space as the thioxoimidazolidinone pharmacophore, thereby mimicking the network of hydrogen bond donors/acceptors required for activity.8 This approach is illustrated in Figure 2 and highlights key objectives which include removal of the Michael acceptor and elimination of isomeric mixtures.

For the current study we elected to retain the isobenzofurane C-subunit of 2 as this class is generally more potent than the corresponding isoinolinones 8 which were originally introduced on the assumption they would be less susceptible to hydrolysis. However subsequent stability studies showed no clear advantage with both the isoinolinones and isobenzofuranones showing variable half-lives in vitro (microsome stability) and in vivo.11 Parent compound 6 was employed as a start point for the replacement of the 2-thioxoimidazolidin-4-one A-subunit, but due to its lack of potency (Jurkat IC₅₀ >20 μM; Table 1) a series of simple substituted phenyl based derivatives were designed and investigated.

Compounds 7–15 (Table 1) follow an independent 2, 3, 4 substitution pattern on a benzene A-ring with derivative 15 (4-OMe) demonstrating activity against isolated perforin protein in our Jurkat assay (IC₅₀ = 9.36 μM). The remaining methoxy isomers (13, 14) showed no detectable inhibition, suggesting that substitution para to the connecting thiophene B-subunit is essential for activity. Prompted by this encouraging result and with derivative 15 providing an H-bond acceptor moiety, we looked to introduce an NH₂, OH and CN group at positions 3 and 4 accordingly. Isomers 16 (3-NH₂) and 17 (4-NH₂) both have comparable activity (IC₅₀ = 13.97 and 11.82 μM, respectively), showing that substitution at the 3-position is tolerated in this particular case. This is confirmed further by 3-hydroxyl-containing compound 18 in which a 2-fold increase in potency is achieved (IC₅₀ = 5.89 μM). Looking at 3-CN derivative 20 in comparison, activity is lost altogether suggesting that an H-bond donor may be required in this position for activity. In contrast, 4-OH containing compound 19 loses all activity whilst 4-CN containing compound 21 (IC₅₀ = 6.87 μM) retains potency similar to that of 18. This reinforces the argument in that an H-bond acceptor at position 4 is more favourable and perhaps necessary for activity to exist in this series (15 and 21 vs 19).

Compounds 22 and 23 were designed with a carboxamide moiety installed at positions 3 and 4, respectively. Substitution at position 3 was well tolerated giving rise to an IC₅₀ = 2.97 μM—2-fold greater than our previously most active compound 18. However when the primary amide is moved to position 4, a dramatic increase in the ability to inhibit perforin lytic activity is seen. Benzene-4-carboxamide 23 has a potency approximately 4-fold greater than lead thioxoimidazolidine 2 (Figs. 1 and 2) against isolated perforin (0.18 μM vs 0.78 μM, respectively). Furthermore, the corresponding pyridine-4-carboxamide 67 is also one of our most potent compounds (IC₅₀ = 0.92 μM). Although at the outset a decision was made to retain the more potent isobenzofuranone C-subunit, for completeness the analogous isoinolin-1-one (23a)
Table 1
Inhibitory activities of isobenzofuran-1(3H)-ones with various A-subunits

| Number | Scheme | X | Inhibition of Jurkat cell lysis IC₅₀ᵇ (µM) |
|--------|--------|---|------------------------------------------|
| 6      | 1      | Ph | >20                                      |
| 7      | 1      | 2-Me Ph | >20                                      |
| 8      | 1      | 3-Me Ph | >20                                      |
| 9a     | 1      | 4-Me Ph | >20                                      |
| 10     | 1      | 2-Cl Ph | >20                                      |
| 11     | 1      | 3-Cl Ph | >20                                      |
| 12     | 1      | 4-Cl Ph | >20                                      |
| 13     | 1      | 2-Ome Ph | >20                                      |
| 14     | 1      | 3-Me Ph | >20                                      |
| 15     | 1      | 4-Ome Ph | >20                                      |
| 16     | 1      | 3-NH₂ Ph | 13.97                                   |
| 17     | 1      | 4-NH₂ Ph | 11.82                                   |
| 18     | 1      | 3-OH Ph | 5.89                                     |
| 19     | 1      | 4-NH₂ Ph | >20                                      |
| 20     | 1      | 3-CN Ph | 20                                       |
| 21     | 1      | 4-CN Ph | 6.87                                     |
| 22     | 1      | 3-CONH₂ Ph | 2.97                                   |
| 23     | 1      | 4-CONH₂ Ph | 0.18                                   |
| 23a    | 1      | 4-CONH Ph | 6.04                                    |
| 29     | 2      | 4-CONMe₂ Ph | 10.7                                    |
| 30     | 2      | 4-CONMe₂ Ph | >20                                    |
| 31ᵃ     | 1      | 4-CONH(CH₂)₃NMe₂ Ph | >20                                   |
| 32     | 1      | 4-CONH(CH₂)₃NMe₂ Ph | >20                                   |
| 39     | 3      | 4-COOEt Ph | 10.97                                   |
| 40     | 3      | 3-OH, 4-COOEt Ph | >20                                   |
| 46     | 3      | 3-NH₂, 4-COOH Ph | 5.84                                    |
| 48     | 3      | 3-OH, 4-CONH₂ Ph | 0.67                                    |
| 49ᵇ     | 3      | 3-CONH(CH₂)₂NMMe₂ Ph | 12.75                                   |
| 50ᵇ     | 3      | 3-OH, 4-CONH(CH₂)₂NMMe₂ Ph | >20                                   |
| 51     | 3      | 3-OH, 4-CONHCH₂CH₂OH Ph | 3.31                                   |
| 52     | 3      | 3-OH, 4-CONHCH₂OHCH₂ Ph | >20                                   |
| 53     | 3      | 3-OH, 4-CONH₂CONH₂ Ph | 2.65                                    |
| 54     | 3      | 3-OH, 4-CONH₂CONH₂ Ph | 7.12                                    |
| 55     | 3      | 3-NH₂, 4-CONH₂ Ph | 1.20                                    |
| 56ᵇ     | 3      | 3-NH₂, 4-CONH(CH₂)₂NMMe₂ Ph | >20                                   |
| 57ᵇ     | 3      | 3-NH₂, 4-CONH(CH₂)₂NMMe₂ Ph | >20                                   |
| 58     | 1      | 4-SMe Ph | >20                                      |
| 59     | 1      | 4-Ch₂OH Ph | 0.90                                     |
| 60     | 1      | 4-SO₂Me Ph | >20                                      |
| 61     | 1      | 3-SO₂NH₂ Ph | 4.85                                    |
| 62     | 1      | 4-SO₂Ntert-Bu Ph | 6.09                                    |
| 63     | 1      | 4-NHSO₂Me Ph | 1.09                                    |
| 64     | 7      | 4-CONH₂, 3-pyridyl Ph | 0.92                                    |
| 70     | 1      | 4-Pyridyl | 4.53                                    |
| 71     | 1      | 5-Isobenzofuran-1(3H)-one | 6.24                                    |

ᵃ Structure of the A-subunit.
ᵇ Testing was carried out over a range of doses, with the IC₅₀ being equal to the concentration at which 50% inhibition of the lysis of Jurkat cells by perforin was observed, as measured by ⁵¹Cr release. Values are the average of at least two independent IC₅₀ determinations.
ᶜ Isobenzofuran-1(3H)-one C-subunit replaced with an isoinolin-1-one.
ᵈ Isobenzofuran-1(3H)-one C-subunit replaced with an 2-methylisooindolin-1-one.
ᵉ Detailed structures for compounds, 31, 49, 50, 56 and 57 can be found in the Supplementary information (Experimental Section).

and 2-methylisooindolin-1-one (23b) derivatives were also prepared. As expected this modification resulted in a loss of activity. Conversion of the primary amide of 23 to a secondary derivative (29) results in a loss of efficacy, while the presence of a tertiary amide (30, 31) abolishes activity completely. Esters 39 showed limited activity (10.97 µM), while compounds 40, 46, 48 and 55 were designed in an effort to combine the outstanding potency of 23 with the preferred 3-OH of 18 or 3-NH₂ of 16, thereby introducing an H-bond donor at the 3-position and adding an ionisable centre to assist solubility. While this approach did generate potent (and slightly more soluble—see Supplementary data) compounds, none were an improvement on 23. In an extension of this strategy, solubilising sidechains were introduced in examples 32, 48–54 and 56, 57. Disappointingly, compounds with strongly or weakly basic sidechains (32, 49, 50, 56, 57) displayed poor activity or were inactive, while neutral compounds 51 and 53 showed only moderate activity (IC₅₀ 3.31 and 2.65 µM), respectively.

In order to further increase diversity and enrich the SAR of the diarylthiophene series, a range of commercially available boronates were deployed in Suzuki reactions with key intermediate 5 (Scheme 1), generating analogues 58–63. Results were mixed, with the methyl alcohol 59 and methyl sulfonamide 63 (IC₅₀ 0.92 and 1.09 µM, respectively) the best of this set. Finally, given similarities in SAR between the A- and C-subunits the symmetrical analogue 71 was synthesised but proved only moderately potent as well as extremely insoluble.

Biological activity and stability: The five most potent inhibitors of isolated recombinant perforin (23, 48, 59, 63, and 67 from Table 1) were then subjected to more advanced assessment. Preliminary stability studies were carried out by incubation in human or mouse plasma at 37 °C with the percentage parent remaining measured at 24 h (Table 2). All five inhibitors were significantly more stable in mouse plasma compared to human. This result is not unexpected, as the antitumor agent camptothecin which also contains a lactone moiety has been shown to co-exist in both the closed and ring-opened form and that this equilibrium is distinctly different for human and mouse plasma. In mouse plasma the ratio of open to closed is 50:50, while in human plasma this shifts to 90:10 due to the strong affinity of human serum albumin for the ring-opened form.²⁰ It is likely that the same phenomenon is operating in the present case and while not necessarily an issue for mouse studies, will need to be addressed in future work.

The compounds were then evaluated for their ability to block the effect of perforin delivered by whole KHYG-1 NK cells (Table 2). Employing whole NK cells is a more rigorous model of conditions in vivo than the use of isolated recombinant protein since effector cell identification of the labelled target cell and formation of an immune synapse are required for perforin release. Any putative inhibitor must also possess the ability to access perforin released into the synaptic cleft. After co-incubation of inhibitor with KHYG-1 NK cells in medium for 30 min at room temperature, ³¹Cr-labelled K562 leukaemia target cells were added and cell lysis evaluated after 4 h incubation at 37 °C by measuring ³¹Cr release. The viability of the NK cells in the presence of inhibitor was also assessed 24 h. later to confirm that any inhibitory activity exhibited by the compounds was due to blocking the action of perforin and not nonspecific killing of the effector cells. Viable and dead cells were counted and percent viability calculated based on total cell count (for further details see Supplementary data). Of the five compounds selected for further testing, three showed poor or no activity; the 3-OH, 4-CONH₂ compound 48, 4-methyl alcohol 59 and methanesulfonamide 63. Gratifyingly, the most potent compound against isolated perforin (23) showed comparable activity (60% inhibition of lytic activity at 20 µM concentration) to many of the active compounds contained in our previous report on the thiooximidazolidinone-based series. In addition, pyridinecarbox-
amidine 67 was essentially equivalent to our previously most potent in vivo candidates (80% inhibition of lysis). Most importantly, this activity was observed in the absence of toxicity toward the effector cells (90% and 82% viability, respectively). For the purposes of this work, compounds are classified as toxic if NK cell viability falls below 70%. This is due to the importance of CTLs and NK cells in the overall immunological response which means that it is essential that any pharmacological intervention allows rapid recovery of these cytotoxic effector cells for normal immunological function.

In summary, a new series of benzene- and pyridine-carboxamides have been designed as isosteric replacements for our previous thioxoimidazolidinone-based series. Several of these compounds showed improved inhibition of perforin-mediated lysis, with benzene-carboxamide 23 demonstrating a 4-fold increase (0.18 μM) over the corresponding thioxoimidazolidinone inhibitor 2 (0.78 μM). Compound 23 and its pyridine analogue 67 also showed
excellent activity against the lytic action of whole human NK cells, at a level comparable to our previously identified in vivo candidates. Use of an isostere in place of the thioxoimidazolidinone A-subunit has also enabled removal of a potentially reactive Michael acceptor moiety and elimination of E- and Z-isomeric mixtures. Finally, these compounds do not appear to possess the marginal toxicity associated with many of the thioxoimidazolidinones.

We can conclude that we have identified viable replacements for the thioxoimidazolidinone subunit while improving potency. We have also overcome several key concerns in relation to this historic series and now seek to expand on these findings to identify a suitable in vivo candidate. The mechanism of action of these inhibitors is still under investigation, however given our previous conclusion that this class of compounds target perforin after its release into the immune synapse it is most likely that inhibition occurs by either (1) blocking the monomers from association with the target cell membrane, (2) counteracting the oligomerisation process that forms the pre-pore, or (3) by blocking of the conformational changes required for pore formation. This work is currently ongoing and further developments will be reported in the future.

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at http://dx.doi.org/10.1016/j.bmcl.2015.12.003.

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