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Metal stopping reagents facilitate discontinuous activity assays of the de novo purine biosynthesis enzyme PurE

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Abstract

The conversion of 5-aminoimidazole ribonucleotide (AIR) to 4-carboxy-AIR (CAIR) represents an unusual divergence in purine biosynthesis: microbes and nonmetazoan eukaryotes use class I PurEs while animals use class II PurEs. Class I PurEs are therefore a potential antimicrobial target; however, no enzyme activity assay is suitable for high throughput screening (HTS). Here we report a simple chemical quench that fixes the PurE substrate/product ratio for 24 h, as assessed by the Bratton-Marshall assay (BMA) for diazotizable amines. The ZnSO₄ stopping reagent is proposed to chelate CAIR, enabling delayed analysis of this acid-labile product by BMA or other HTS methods.

Keywords: purine biosynthesis, aminoimidazole, substrate depletion, chelation

1 De novo purine biosynthesis harbors an unusual divergence in a ubiqui-

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2 uitous biosynthetic pathway [1]. In most microbes and nonmetazoan eu-
3 karyotes, a class I PurE [PurE1, N^5 -carboxyaminoimidazole ribonucleotide
4 (NCAIR) mutase, EC 5.4.99.18] reversibly converts NCAIR to 4-carboxy-5-
5 aminoimidazole ribonucleotide (CAIR) (Fig. 1) [2]. Animals produce CAIR
6 by a different route, using a class II PurE (PurE2, 5-aminoimidazole ribonu-
7 cleotide (AIR) carboxylase, EC 4.1.1.21) [3, 4, 5]. Microbial pathogenesis
8 relies upon purine biosynthesis [6], particularly *purE1*, which is therefore a
9 promising but unexploited antibiotic target [7, 8]. Inhibitors selective for
10 PurE1 could have antibacterial, antifungal, or herbicidal activity [9]. In this
11 work, we describe a functional assay for PurE that is compatible with in-
12 hibitor identification by high-throughput screening (HTS) methods.

13 Currently available PurE functional assays are not amenable to HTS
14 methods. A continuous ultraviolet (UV) assay that monitors the absorbance
15 decrease due to CAIR decarboxylation [10, 11, 2] is subject to spectral in-
16 terference by many candidate inhibitors. CAIR synthesis assays are compli-
17 cated by the acid lability of the NCAIR substrate [12, 13, 2], the need for
18 multiple coupling enzymes [14, 9], and, for discontinuous assays detecting in-
19 organic phosphate production, handling of malachite green reagent [15] and
20 a background due to formation of phosphate by the coupling enzymes. The
21 Bratton-Marshall assay (BMA) [16] was used in early studies of PurE2 [3],
22 but seldom thereafter. The BMA yields intensely colored AIR and CAIR
23 derivatives, AIR-BM and CAIR-BM (Fig. 1), that possess characteristic vis-
24 ible absorbance spectra [17], are relatively stable, and might be suitable for
25 HTS methods. The BMA, however, requires multiple timed reagent additions
26 that would be difficult to implement during parallel screens. An alternative

27 would be to stop the PurE reaction prior to BMA workup using a reagent
28 that does not alter product yields.

29 The first task was to determine an extinction coefficient for CAIR-BM at
30 500 nm, a goal that requires pure CAIR. A new CAIR synthesis, which uses
31 LiOH to saponify aminoimidazole-4-carboxamide ribonucleotide (AICAR)
32 [18] and thereby facilitates salt removal (see Experimental Procedures in the
33 Supplementary material), furnished CAIR in high yield (85%) and purity
34 (99%). No signal corresponding to AIR H2 was observed in ^1H NMR spectra
35 (Fig. S1 in the Supplementary material). Mass spectrometry (MS) analy-
36 sis of BMA reaction mixtures revealed only the expected molecular ions for
37 single BM adducts and ions present in control mixtures (Table S1 in the Sup-
38 plementary material). HPLC analysis of CAIR subjected to BMA workup
39 showed 97% CAIR-BM / 3% AIR-BM (Fig. S2 in the Supplementary mate-
40 rial); a small amount of AIR evidently forms during the BMA acid quench
41 step. CAIR-BM appears to consist of a mixture of species that cannot be
42 resolved by HPLC but that possess the same molecular formula. An isobaric
43 mixture might also account for the complex shape of the CAIR-BM visible
44 spectrum (Fig. S3 in the Supplementary material). Aliquots of a solution
45 containing pure CAIR were (1) used to determine an accurate CAIR concen-
46 tration, by complete enzymatic conversion to AIR followed by BMA analysis,
47 and (2) used to obtain a spectrum of CAIR-BM, which gave $\epsilon_{500}^{\text{CAIR-BM}} = 11\,200$
48 $\text{M}^{-1} \text{cm}^{-1}$ (for comparison, $\epsilon_{500}^{\text{AIR-BM}} = 24\,800 \text{M}^{-1} \text{cm}^{-1}$ [17]).

49 The second task was to identify an enzyme quenching method that is
50 compatible with BMA workup and product analysis. CAIR and analogues
51 form stable complexes in which aminoimidazole N3 and O7 atoms (Fig. 1)

52 coordinate a transition metal ion [19]. (CAIR)_n-metal ($n = 2$ or 3) complexes
53 (Fig. S4 in the Supplementary material) do not bind to or inhibit chicken
54 PurE2 in the presence of excess free CAIR [20, 21, 22]. Mg²⁺ inhibits PurE2-
55 mediated CAIR decarboxylation but not AIR carboxylation, consistent with
56 the formation of a low-affinity metal-CAIR complex [10, 4]. To determine if
57 transition metals can rapidly and reversibly form CAIR-metal complexes, a
58 ten-fold molar excess (relative to the initial CAIR concentration) of NiSO₄ or
59 ZnSO₄ was added to UV-monitored CAIR decarboxylation reaction mixtures.
60 CAIR decarboxylation stopped immediately, but could be restarted by the
61 addition of EDTA (Fig. S5 in the Supplementary material). EDTA had no
62 effect on BMA workup; the initial acidification step apparently dissociates
63 CAIR complexes formed with either Ni²⁺ or Zn²⁺.

64 Zn²⁺ appeared to form a CAIR complex with higher kinetic stability than
65 Ni²⁺, and to pose fewer concerns with regard to toxicity, spectral overlap
66 with (C)AIR-BM, and redox chemistry. A CAIR decarboxylation reaction
67 mixture incubated at 30 °C was stopped at ~30% completion by the addition
68 of ~3 molar equivalents of ZnSO₄ (Fig. 2). Equivalent yields of CAIR-BM
69 were obtained if the BMA workup was delayed by up to 4 h. In contrast,
70 Ni²⁺-stopped reactions lost CAIR at the rate of about 10% per h at 30
71 °C (data not shown). A saturation curve employing ZnSO₄ quenching gave
72 kinetic constants similar to the standard continuous UV assay (Fig. S6 in
73 the Supplementary material). Indistinguishable velocities were obtained at
74 40 μM CAIR for a quenched reaction worked up promptly or after a 24 h
75 delay. Therefore ZnSO₄ is a suitable stopping reagent for PurE reaction
76 mixtures.

77 Metal stopping reagents are proposed to work by rapid substrate de-
78 pletion, an uncommon mode of enzyme inhibition (e.g., vancomycin [24])
79 [25]. Immobilized metal affinity chromatography (IMAC) was used to test if
80 CAIR binds to Ni^{2+} , immobilized on a nitrilotriacetic acid (NTA) column.
81 An equimolar mixture of CAIR and AIR was applied to the NiNTA column.
82 The first column fraction contained pure AIR, consistent with selective re-
83 tention of CAIR on the column as a CAIR- Ni^{2+} complex (Figs. S7 and S8 in
84 the Supplementary material). Subsequent fractions contained CAIR, indicat-
85 ing that the Ni^{2+} -CAIR interaction was reversible (data not shown). While
86 these results do not exclude the possibility that PurE is inhibited by direct
87 protein-metal interactions, they indicate that the rapid formation of a stable
88 CAIR-transition metal complex is sufficient to account for ZnSO_4 -mediated
89 arrest of CAIR decarboxylation.

90 In summary, we have identified a PurE reaction quencher that facilitates
91 HTS using the BMA as a detection method. PurE-mediated CAIR decar-
92 boxylation stopped by a single liquid addition, of a ZnSO_4 solution, allows
93 delayed BMA workup. The BMA is compatible with candidate inhibitors
94 that absorb UV light, a common practical concern for detecting initial hits,
95 which often have low potency and therefore require high compound concen-
96 trations that cause significant spectral interference. BMA-detected PurE
97 screens require no other enzymes and can be used to screen candidate in-
98 hibitors that do not contain diazotizable amines or strong metal chelators.
99 The metal stopping reagent could be adapted for use with other detection
100 methods or in assays of 5-aminoimidazole-4-*N*-succinylcarboxamide ribonu-
101 cleotide (SAICAR) synthetase (PurC, EC 6.3.2.6), the other enzyme known

102 to use CAIR as a substrate. SAICAR has recently been shown to activate
103 cancer metabolism via pyruvate kinase isoform M2 [26].

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214 **Figure Captions**

215 **Figure 1:** PurE1 reaction and CAIR/AIR detection by Bratton-Marshall
216 assay (BMA). NCAIR decarboxylates to AIR (half-life of 0.9 min at pH 7.8
217 and 30 °C) during the enzyme assay [2], an acid-dependent process that goes
218 to completion during the BMA. R5P = ribosyl 5'-phosphate, R = H (AIR-
219 BM) or COO⁻ (CAIR-BM). CAIR aminoimidazole atoms are numbered as
220 in PDB residue type C2R. CAIR atoms proposed to coordinate transition
221 metals are indicated with unfilled arrows.

222 **Figure 2:** Zn²⁺ ions stop CAIR decarboxylation. Reaction mixtures (1.26
223 mL final volume) at 30 °C contained 50 mM Tris•HCl, pH 8.0, 42 μM CAIR,
224 and 0.13 μg *Acetobacter aceti* PurE1 (0.005 units) [23]. At 5 min, ZnSO₄
225 (0.1 mM) was added to one reaction. Aliquots (0.2 mL) were removed prior
226 to the addition of enzyme (zero time point) and at indicated intervals, and
227 immediately added to a 1.7 mL centrifuge tube containing a freshly prepared
228 mixture of 20% (w/v) trichloroacetic acid / 1.33 M potassium phosphate,
229 pH 1.4, (0.1 mL) and 0.1% (w/v) sodium nitrite (0.1 mL). After 3 min, 0.5%
230 (w/v) ammonium sulfamate (0.1 mL) was added to destroy excess nitrite.
231 After an additional min, Bratton-Marshall reagent [0.1 mL of 0.1% (w/v)
232 *N*-(1-naphthyl)ethylenediamine dihydrochloride] was added. After 10 min,
233 the solution (0.6 mL) was centrifuged (16 000*g*, 30 s) to eliminate bubbles
234 and the absorbance at 500 nm (A_{500}) was recorded.

235 CAIR consumption was quantitated using a differential extinction coef-
236 ficient for CAIR to AIR conversion, $\Delta\epsilon_{500} = \epsilon_{500}^{\text{AIR-BM}} - \epsilon_{500}^{\text{CAIR-BM}} = 13\,600 \pm$
237 $1600 \text{ M}^{-1} \text{ cm}^{-1}$. [CAIR] values at time t were determined using ($A_{500}^{\text{AIR-BM}} -$

238 $A_{500}^t)/\Delta\epsilon_{500}$, where $A_{500}^{\text{AIR-BM}}$ was from a control reaction in which CAIR was
239 completely converted to AIR.

240 The symbols and error bars represent the mean and standard deviation,
241 respectively, for at least three independent replicates of ZnSO_4 -quenched
242 (filled circles) and -unquenched control (open circles) reactions. Concentra-
243 tions determined at $t > 5$ min were significantly different in quenched and
244 control reaction mixtures [one-tailed t -test, P -value < 0.01 (**) or < 0.001
245 (***)]. The slope of the line connecting the first two points corresponds to
246 0.003 units of enzyme activity, comparable to the 0.004 units expected under
247 the reaction conditions [23].