SYNTHESIS AND RESOLUTION OF 7-FLUOROTRYPTOPHANS¹

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Abstract: 7-Fluoro-DL-tryptophan (1a), 4,7-difluoro-DL-tryptophan (1b) and 5,7-difluoro-DL-tryptophan (1c) were prepared by Fischer indole cyclization. The resolution was achieved by treatment of the N-trifluoroacetyl derivatives with carboxypeptidase A. All 7-fluorotryptophans are slow substrates for tryptophan indole-lyase from E. coli.

Fluorinated analogues of amino acids are useful to study amino acid reactions and metabolism.² Recently, there has been an increasing interest in the incorporation of fluoroamino acids in large biomolecules.³ F-Phe-, F-Trp-, and F-Tyr-labeled proteins have been used in ¹⁹F NMR studies to characterize fuctionally significant conformational changes.⁴ Among fluorotryptophans, 4-, 5- and 6-fluorotryptophan are commercially available, and used extensively.^{2c,3b,5} 4, 5, 6, 7-Tetrafluorotryptophan was synthesized and tested for enzyme inhibition.⁶ However, there is only one report for preparation of 1a and 1c, which were not resolved,^{6e} and the preparation of 1b has not been previously reported. We report here the synthesis of 7-fluorotryptophans 1a-c, their resolution, and their reactions with tryptophan indole-lyase (tryptophanase) from *E. coli*.

Fluoroanilines were diazotized, reduced with SnCl₂, and the resulting HCl salts were neutralized with 4N NaOH solution to give fluorophenylhydrazines 2a-c in 67-85% yields.⁷ After the reaction of diethyl acetamidomalonate with acrolein and sodium methoxide in benzene,⁸ addition of the fluorophenylhydrazine gave the crystalline hydrazones 3a-c. (85-90%) The predominant stereochemistry of the hydrazones across the imine double bond was determined to be E, based on the results of the proton NMR.⁹ Reflux of 3a-c with dilute sulfuric acid (5-10%) for 5 hours afforded

indoles 4a-c.¹⁰ The yields were dependent on the number and position of fluorines, and were 38%, 36% and 12% for 4a, 4c and 4b, respectively. Saponification and decarboxylation of 4a-c were achieved by refluxing with 1.2 equivalent NaOH in aqueous dioxane (1:1).¹¹ The resulting N-acetyltryptophans 5a-c were hydrolyzed

with 4 equivalents of NaOH to give 1a-c.¹² 1a-c were converted to the N-trifluoro-acetyl derivatives¹³ and were treated with carboxypeptidase A to obtain the 7a-c and 8a-c.¹⁴ The N-TFA-D-7-fluorotryptophans were treated with 1M aqueous piperidine to give the D-7-fluorotryptophans 9a-c. The optical purity of each enantiomer was measured by HPLC using a chiral column, and was greater than 99% e.e.¹⁵ We have examined the kinetics of the reaction of Escherichia coli tryptophan indole-lyase with compounds 7a-c.¹⁶ All of them are slower substrates than L-tryptophan.

| Table | 1. | Kinetic | data | for | 7a-c | with | tryptophan | indole-lyase |
|-------|----|---------|------|-----|------|------|------------|--------------|
| | | | | | | | | |

| Compound | $K_m(mM)$ | $V_{max}(rel.)$ | $k_{cat}/K_m(rel.)$ |
|--------------|-----------|-----------------|---------------------|
| L-tryptophan | 0.26 | 100% | 100% |
| 7 a | 0.64 | 29% | 14% |
| 7 b | 1.7 | 42% | 6% |
| 7 c | 0.49 | 58% | 30% |

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References and Notes

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- Each compound was characterized by ¹H, ¹³C and ¹⁹F NMR, and elemental 12. analysis. 1a: mp 252-254 °C(lit. 254-256 °C); ¹H NMR (Methanol-D4, 300Mhz, ppm) 7.49 (d, 1H, J=8.0, C4-H), 7.23(s, 1H, C2-H), 6.98(td, 1H, J=4.7, 7.9Hz, C5-H), 6.84(dd, 1H, J=7.9, 11.5Hz, C6-H), 3.84(dd, 1H, J=4.1, 9.3Hz, αC-H), 3.49(dd, 1H, J=4.1, 15.3Hz, βC-H), 3.15(dd, 1H, J=9.3, 15.2Hz, βC-H) 13 C NMR (Methanol-D₄, 62.9 Mhz, decoupler on, ppm) 174.2(COO), 151.2(d, J_{C-F}=-242.9Hz, C7), 115.4(d, J_{C-F} F=6.1Hz, C3a),126.3(d, J_{C-F}=13Hz, C7a), 126.2(C2), 120.4(d, J_{C-F}=6.2Hz, C5), 114.5(d, $J_{C-F}=3.3Hz$, C4), 110.7(C3), 107.3(d, $J_{C-F}=16.4Hz$, C6), 56.6(α C), 28.4(β C), ¹⁹F NMR(Methanol-D₄, 282.4Mhz, ppm) -135.2(dd, J=4.7, 11.5Hz). Analysis. Calc.: C 59.46% H 4.95% N 12.6%; Found: C 59.36%, N 5.01%, H 12.58% **1b**: mp 264-265 °C; ¹H NMR (Methanol-D₄, 300Mhz, ppm) 7.24(s, 1H, C₂-H), 6.80(ddd, 1H, J=3.5, 8.5, 10.4Hz, C5-H)*, 6.67(ddd, 1H, J=3.1, 8.5, 10.5Hz, C6-H)*, 4.22(dd, 1H, J=5.0, 9.3Hz, αC-H), 3.60(dd, 1H, J=5.0, 15.0Hz, βC-H), 3.25(dd, 1H, J=9.3, 15.0Hz, βC-H) ¹³C NMR (Methanol-D₄, 75Mhz, decoupler on, ppm) 171.4 (COO), 154.1(dd, J_C. F=1.9, -238.4Hz, C4), 147.6(dd, $J_{C-F}=2.7$, -239.2Hz, C7), 128.6(dd, $J_{C-F}=11.6$, 16.6Hz, C7a), 127.2(C2), 119.6(dd, J_{C-F}=22.5, 12.5Hz, C3a), 108.1(C3), 107.3(dd, J_{C-F}=22.5, 12.5Hz, C3a), 107.3(dd, J_{C-F}=22.5, 12 F=8.9, 19.6Hz, C5)*, 104.6(dd, $J_{C-F}=7.1$, 23.0Hz, C6)*, 56.0 (α C), 28.6(β C), ¹⁹F NMR(Methanol-D4, 282.4Mhz, ppm) -129.8(ddd J=3.5, 10.5, 22.5, F4) -139.6(ddd, J=3.1, 10.4, 22.5, F7), Analysis. Calc.: C 55% H 4.2% N 11.66%; Found: C 54.92%, H 4.2%, N 11.62% * interchangable 1c: mp 249-250 °C; ¹H NMR (Methanol-D₄, 300Mhz, ppm) 7.30 (s, 1H, C2-H), 7.26(dd, 1H, J=2.2, 9.4Hz, C4-H), 6.73(ddd, 1H, J=2.2, 9.6, 11.1Hz, C6-H), 3.82(dd, 1H, J=4.3, 8.7Hz, α C-H), 3.40(dd, 1H, J=4.3, 15.3Hz, β C-H), 3.15(dd, 1H, J=8.7, 15.3Hz, β C-H) ¹³C NMR (Methanol-D₄, 62.9Mhz, decoupler on, ppm) 174.1(COO), 158.1(dd, J_{C-F}=9.7, -235.0Hz, C5), 150.2(dd, J_{C-F} F=14.3, -246.1Hz, C7), 131.4(dd, $J_{C-F}=6.7$, 11.2Hz, C3a), 128.0(C2), 122.9(d, $J_{C-F}=6.7$, 12.2Hz, C3a), 128.0(C2), 128.0(13.5Hz, C7a), 110.9(C3), 100.4(dd, J_{C-F}=3.7, 23.8Hz, C4), 97.4(dd, J_{C-F}=20.9, 30.7Hz, C6), 56.4(αC), 28.1(βC), ¹⁹F NMR(Methanol-D₄, 282.4Mhz, ppm) -122.1 (ddd, J=2.1, 9.4, 9.6Hz, F5) -131.3(dd, J=2.1, 11.1Hz, F7) Analysis. Calc.: C 55% H 4.2% N 11.66%; Found: C 54.94%, H 4.23%, N 11.57%.
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- 15. Column: Cu-proline. Eluent: 10mM CuSO₄ Detector: 270nm L-isomer elutes last. 7a; $[\alpha]^{25}_{D}$ = -21.8 (c=0.7, DMSO) 7b; $[\alpha]^{25}_{D}$ = -44.1(c=0.74, formamide) 7c; $[\alpha]^{25}_{D}$ = -27.5 (c=1, methanol) 9a; $[\alpha]^{25}_{D}$ = +23.1(c=0.73 DMSO) 9b; $[\alpha]^{25}_{D}$ = +45.9 (c=0.73, formamide) 9c; $[\alpha]^{25}_{D}$ =+28.1 (c=1, methanol)
- 16. Tryptophan indole-lyase was purified from E. coli JM101 containg the tnaA gene on plasmid pMD6 (Phillips, R. S. and Gollnick, P. D., J. Bio. Chem., 1989, 264, 10627). Catalytic turnover with L-tryptophan and L-7-fluoro-tryptophans was measured by coupled assay with lactate dehydrogenase and NADH at 340nm. (Torino, Y., and Snell, E. E., Methods Enzymol., 1970, 17A, 439).