### **Electronic Supplementary Information for**

## Chiral Discrimination of α-Amino Acids with a C<sub>2</sub>-Symmetric Homoditopic Receptor

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## CONTENT

1.	General Methods
2.	Synthesis of receptor (S)-1
3.	Preparation of various amino acid tetrabutylammonium carboxylates
4.	Figure S1. Proton-coupled ${}^{19}F$ NMR spectra of (S)-1 with alanine
	tetrabutylammonium carboxylate
5.	Figure S2. Proton coupled ${}^{19}F$ NMR spectra of (S)-1 with serine
	tetrabutylammonium carboxylate
6.	Figure S3. Proton coupled <sup>19</sup> F NMR spectra of (S)-1 with tryptophan
	tetrabutylammonium carboxylate
7.	Figure S4. Proton coupled $^{19}$ F NMR spectra of (S)-1 with phenylglycine
	tetrabutylammonium carboxylate
8.	Figure S5. Proton coupled $^{19}$ F NMR spectra of (S)-1 with phenylalanine
	tetrabutylammonium carboxylate
9.	Figure S6. Proton coupled $^{19}$ F NMR spectra of (S)-1 with methionine
	tetrabutylammonium carboxylate
10.	Figure S7. Proton coupled ${}^{19}F$ NMR spectra of (S)-1 with value
	tetrabutylammonium carboxylate
11.	Figure S8. Proton coupled <sup>19</sup> F NMR spectra of (S)-1 with leucine
	tetrabutylammonium carboxylate
12.	Figure S9. Proton coupled $^{19}$ F NMR spectra of (S)-1 with tetrabutylammonium
	acetate, 1,2-diaminopropane and glycine tetrabutylammonium
	carboxylate
13.	Figure S10. Partial <sup>1</sup> H NMR spectra of (S)-1–L-alanine tetrabutylammonium

	carboxylate adduct and receptor (S)-1-D-alanine tetrabutylammonium
	carboxylate adduct
14.	Figure S11. Partial <sup>1</sup> H NMR spectra of receptor (S)-1–L-alanine
	tetrabutylammonium carboxylate adduct and receptor (S)-1-D-alanine
	tetrabutylammonium carboxylate adduct
15.	Figure S12. <sup>13</sup> C NMR spectra of (S)-1–L-alanine tetrabutylammonium
	carboxylate adduct and (S)-1-D-alanine tetrabutylammonium carboxylate
	adduct
16.	<b>Figure S13</b> . Partial <sup>1</sup> H- <sup>1</sup> H COSY spectrum of ( <i>S</i> )-1S19
17.	Figure S14. Isothermal titration calorimetry data of (S)-1 with L-alanine and D-
	alanine tetrabutylammonium carboxylates
18.	Figure S15. Reversibility of alanine binding to ( <i>S</i> )-1
19.	<sup>1</sup> H and <sup>13</sup> C NMR spectra of compounds ( <i>S</i> )-1

### **General Methods**

All moisture or air sensitive experiments were performed under a positive pressure of argon in oven dried glassware equipped with a rubber septum inlet. Solvents and liquid reagents were transferred by Argon flushed syringes or cannula. Reaction mixtures were stirred with Teflon coated magnetic stir bars. Reaction solvents used were HPLC grade, dried further by passage through a column of appropriate dessicant. Commercial reagents were used as such unless otherwise noted. Analytical thin layer chromatography was performed using Merck 60 F<sub>524</sub> pre-coated silica gel plates. Subsequent to elution, the plates were visualized by ultra-violet light at 254nm. Preparative thin layer chromatography was also performed using Merck 60 F<sub>524</sub> precoated silica gel plates of thickness 1 mm. Flash column chromatography was performed using Merck silica gel 60 (230-400 mesh). Melting points were measured on a capillary melting point apparatus heated by a heating block and are uncorrected. The nuclear magnetic resonance spectra were recorded on an AM-300 Bruker (<sup>1</sup>H, 300 MHz or 500MHz and <sup>13</sup>C, 75 MHz). Chemical shifts are reported in parts per million using the residual solvent peaks as internal reference. The proton coupled <sup>19</sup>F NMR spectra are reported in parts per million (ppm) relative to trifluoroacetic acid ( $\delta$  0.0 ppm) as an external reference. Mass spectral analysis was recorded on JEOL JMS-AX505WA and is reported in units of mass to charge (m/z). HR-MS was performed at the Korea Basic Science Center, Kyungpook National University



#### Synthesis and characterization of receptor (S)-1

Synthesis of receptor (S)-1 was carried out starting from commercially available and optically pure (S)-(-)-2.2'-dihydroxy-1,1'-binaphthyl, according to the scheme above. As all the steps involve known non-racemization processes, we presume that the optical purity of the starting material is transmitted to the final product, receptor (S)-1. A detailed synthesis of (S)-1 and its analogues will be reported in a full account.

Receptor (*S*)-1. (*S*)-2,2'-Diisopropoxy-1,1'-binaphthyl-3,3'-dicarboxylic acid (5), (500 mg, 1.09 mmol) was placed in a dry flask with a stirring bar under argon atmosphere, and freshly distilled  $SOCl_2$  (5 mL) was introduced. The mixture was refluxed for 6 h and excess  $SOCl_2$  was removed under reduced pressure to give the corresponding diacid

chloride, which was used as such without purification.

To a solution of 1-(2-aminophenyl)-2,2,2-trifluoroethanol (6) (835 mg, 4.365 mmol) in anhydrous THF (10 mL) at 0 °C, was added dropwise a solution of the diacid chloride prepared above in anhydrous THF (5 mL). The reaction mixture was stirred for 15 h at room temperature, and the solvent was removed on a rotary evaporator and the resulting residue was purified by column chromatography on silica gel (eluant:  $10 \rightarrow$ 40% EtOAc in hexane) to give (1S)-2,2'-diisopropoxy-N<sup>3</sup>,N<sup>3'</sup>-bis[2-(2,2,2-trifluoro-1hydroxyethyl)phenyl]-1,1'-binaphthyl-3,3'-dicarboxamide (7) as a white solid (777 mg). The product which was a mixture of rotamers and diasteromers was used as such in the next step. To a solution of compound 7 (674 mg, 0.838 mmol) in dichloromethane (10 mL), was added a solution of Dess-Martin periodinane (5.92 mL, 15 wt% in CH<sub>2</sub>Cl<sub>2</sub>, 2.095 mmol) dropwise at 25 °C. The reaction mixture was stirred for 16 h at room temperature, then the insolubles were filtered off and the volatiles were distilled off in vacuo. The residue was dissolved in ethyl acetate (20 mL) and washed with saturated sodium bicarbonate solution, saturated sodium thiosulfate solution, and finally with saturated brine solution. The ethyl acetate solution was then evaporated in vacuo and the residue obtained was subjected to silica gel column chromatography (eluant:  $5 \rightarrow 7\%$ EtOAc in hexane) to afford the product as an off-white solid (490 mg). The product was further purified by preparative TLC (eluant: 15% EtOAc in hexane) to give pure (S)-1 (391 mg, 51 %): Characterization data for (S)-1 : mp 101–103 °C;  $[\alpha]^{22}_{D} = +89.9$  (c 1.5, CHCl<sub>3</sub>); <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>): δ 11.79 (s, 2H), 8.77 (d, J = 8.5 Hz, 2H), 8.69 (s, 2H), 8.04 (d, J = 8.1 Hz, 2H), 7.94 (d, J = 7.8 Hz, 2H), 7.74 (t, J = 7.7 Hz, 2H), 7.49 (t, J = 7.3 Hz, 2H), 7.39 (t, J = 7 Hz, 2H), 7.28 (m, 2H, overlapped with CHCl<sub>3</sub>), 7.24 (m, 2H, overlapped with CHCl<sub>3</sub>), 3.84 (septet, J = 6 Hz, 2H), 0.89 (d, J = 6 Hz, 6H), 0.74 (d,

J = 6 Hz, 6H). <sup>1</sup>H NMR (300 MHz, CD<sub>3</sub>CN):  $\delta$  11.49 (s, 2H), 8.71 (s, 2H), 8.62 (d, J = 8.5 Hz, 2H), 8.18 (dd, J = 8.1,0.36 Hz, 2H), 7.97 (d, J = 8.1 Hz, 2H), 7.84 (ddd, J = 8.1, 7.7, 1.4 Hz, 2H), 7.59 (ddd, J = 10.5, 10.3, 1.1 Hz, 2H), 7.45 (ddd, J = 7.7, 7.6, 1.3 Hz, 2H), 7.39 (ddd, J = 7.7, 7.7, 1 Hz, 2H), 7.25 (d, J = 8.5 Hz, 2H), 3.98 (septet, J = 6.1 Hz, 2H), 0.91 (d, J = 6.1 Hz, 6H), 0.78 (d, J = 6.1 Hz, 6H).<sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>):  $\delta$  181.8 (q, J = 34.5 Hz, <u>-C</u>OCF<sub>3</sub>), 166.5, 151.7, 141.3, 136.6, 135.8, 133.4, 131.2, 130.3, 129.9, 128.9, 126.7, 126.1, 25.9, 123.6, 122.8, 118.8, 116.8 (q, J = 290 Hz, -<u>C</u>F<sub>3</sub>; 114.9 and 111.1 are observed; but other peaks at ~118.8 and 123 are overlapped with other carbons), 78.2, 22.5, 22.1. <sup>19</sup>F NMR (300 MHz, CD<sub>3</sub>CN)  $\delta$  4.8; HRMS calcd for C<sub>44</sub>H<sub>34</sub>F<sub>6</sub>N<sub>2</sub>O<sub>6</sub> [M<sup>+</sup>] 800.2321; found 800.2318.

# Representative procedure for the preparation of various amino acid tetrabutylammonium carboxylates.<sup>2</sup>

To a suspension of L-alanine (250 mg, 2.80 mmol) in water (10 mL) was added tetrabutylammonium hydroxide (1 M in methanol, 2.8 mL, 2.8 mmol). The mixture was stirred and heated to 55 °C for 2 h. The solvents were removed in vacuo and the residue was dissolved in acetonitrile (10 mL) by sonication. The insolubles were filtered off and the filtrate was dried over anhydrous sodium sulfate. The solvent was then evaporated in vacuo and the residue was lyophilized to give the desired amino acid salt as a white solid or semi-solid.

### References

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Figure S1. Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (S)-1 [1.0 mM], (b) (S)-1 [1.0 mM] + L-alanine [3.0 mM], (c) (S)-1 [1.0 mM] + D-alanine [3.0 mM], and (d) (S)-1 [1.0 mM] + DL-alanine [3.0 mM]; taken in CD<sub>3</sub>CN. Alanine is used as its tetrabutylammonium carboxylate salt.



**Figure S2.** Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + L-serine [3.0 mM], (c) (*S*)-1 [1.0 mM] + D-serine [3.0 mM], and (d) (*S*)-1 [1.0 mM] + DL-serine [3.0 mM]; taken in CD<sub>3</sub>CN. Serine is used as its tetrabutylammonium carboxylate salt.



**Figure S3.** Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + L-tryptophan [3.0 mM], (c) (*S*)-1 [1.0 mM] + D-tryptophan [3.0 mM], and (d) (*S*)-1 [1.0 mM] + DL-tryptophan [3.0 mM]; taken in CD<sub>3</sub>CN. Tryptophan is used as its tetrabutylammonium carboxylate salt.



**Figure S4**. Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + L-phenylglycine [3.0 mM] (c) (*S*)-1 [1.0 mM] + D-phenylglycine [3.0 mM], and (d) (*S*)-1 [1.0 mM] + DL-phenylglycine [3.0 mM]; taken in CD<sub>3</sub>CN. Phenylglycine is used as its tetrabutylammonium carboxylate salt.



Figure S5. Proton-coupled <sup>19</sup>F NMR spectra of (a) Receptor (S)-1 [1.0 mM], (b) (S)-1 [1.0 mM] + D-phenylalanine [3.0 mM], (c) (S)-1 [1.0 mM] + DL-phenylalanine [3.0 mM]. and (d) (S)-1 [1.0 mM] + DL-phenylalanine [3.0 mM]; taken in CD<sub>3</sub>CN. Phenylalanine is used as its tetrabutylammonium carboxylate salt.



**Figure S6.** Proton-coupled <sup>19</sup>F NMR spectra of receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + L-methionine [3.0 mM], (c) (*S*)-1 [1.0 mM] + D-methionine [3.0 mM], and (d) (*S*)-1 [1.0 mM] + DL-methionine [3.0 mM]; taken in CD<sub>3</sub>CN. Methionine is used as its tetrabutylammonium carboxylate salt.



**Figure S7.** Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + L-valine [3.0 mM], (c) (*S*)-1 [1.0 mM] + D-valine [3.0 mM], and (d) (*S*)-1 [1.0 mM] + DL-valine [3.0 mM]; taken in CD<sub>3</sub>CN. Valine is used as its tetrabutylammonium carboxylate salt.



**Figure S8.** Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-**1** [1.0 mM], (b) (*S*)-**1** [1.0 mM] + L-leucine [3.0 mM], (c) (*S*)-**1** [1.0 mM] + D-leucine [3.0 mM], and (d) [1.0 mM] + DL-leucine [3.0 mM]; taken in CD<sub>3</sub>CN. Leucine is used as its tetrabutylammonium carboxylate salt.



(a) Receptor 1

**Figure S9.** Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [6.0 mM] +  $Bu_4NAc$  [18.0 mM], (c) (*S*)-1 [6.0 mM] + 1,2-diaminopropane [18.0 mM], and (d) (*S*)-1 [6.0 mM] + glycine [18.0 mM]; taken in CD<sub>3</sub>CN. Glycine is used as its tetrabutylammonium carboxylate salt.



**Figure S10.** Partial <sup>1</sup>H NMR spectra of (a) receptor (*S*)-1 [0.02M], (b) (*S*)-1 [0.02M] + L-alanine [0.02M], and (c) (*S*)-1 [0.02M] + D-alanine [0.02M]; taken in CD<sub>3</sub>CN. Alanine is used as its tetrabutylammonium carboxylate salt.



**Figure S11.** Partial <sup>1</sup>H NMR spectra of (a) receptor (*S*)-**1** [0.02M], (b) (*S*)-**1** [0.02M] + L-alanine [0.02M], and (c) (*S*)-**1** [0.02M] + D-alanine [0.02M]; taken in CD<sub>3</sub>CN. Alanine is used as its tetrabutylammonium carboxylate salt.



**Figure S12.** <sup>13</sup>C NMR spectra of (a) receptor (*S*)-1 [0.02M], (b) (*S*)-1 [0.02M] + Lalanine [0.02M], and (c) (*S*)-1 [0.02M] + D-alanine [0.02M]; taken in CD<sub>3</sub>CN. Alanine is used as its tetrabutylammonium carboxylate salt.

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Figure S13. Partial <sup>1</sup>H-<sup>1</sup>H COSY spectrum of receptor (*S*)-1, taken in CD<sub>3</sub>CN

## Isothermal titration calorimetry (ITC).

The binding affinity and the thermodynamic data were determined by ITC using a isothermal titration calorimeter (Microcal Inc.).

**Procedure:** To a solution of receptor (*S*)-1 in the calorimeter cell, 5.0  $\mu$ L of L- or Dalanine tetrabutylammonium carboxylate was injected 40 times at 25 °C. The dilution effects were corrected by carrying out a separate blank titration. The titration data was analyzed by the built-in curve-fitting Origin software, which gave an apparent binding constant (*K*<sub>ass</sub>) and the standard enthalpy change  $\Delta H^{\circ}$ .

## **Conditions:**

Cell: Receptor (*S*)-1 (0.2 mM) Syringe: L- or D-Alanine tetrabutylammonium carboxylate solution (3.0 mM) Reference power: 26 Temperature: 303 K Stirring rate: 242 rpm

Although the crude binding isotherms have good features as shown in the next figures, we have difficulty in fitting the raw data by nonlinear least squares regression methods implemented in the software. Thus, the nonlinear squares regression fitting procedure gave a large error value ( $\chi^2$  value), particularly in the case of the less stable D-Ala adduct formation. Therefore, the values provided should be taken as ballpark figures. For binding L-alanine tetrabutylammonium carboxylate:  $K_{ass} \approx 1.3 \times 10^7 \text{ M}^{-1}$ , and for binding D-alanine tetrabutylammonium carboxylate:  $K_{ass} \approx 1.1 \times 10^6 \text{ M}^{-1}$ . As shown in the previous reports (references 8 and others reported by Ahn and co-workers for molecular interactions between CATFA-based molecular receptors and carboxylates), the binding process involves favorable enthalpy changes ( $\Delta H = -29$  kcal/mol) and unfavorable entropy changes [ $-T\Delta S = 303 K_{ass} \times (-63 \text{ eu}) = 19 \text{ kcal/mol}$ ] in the case of L-Ala adduct formation.

Even though the fitting procedures gave the ballpark figures for the association constants, the integration data in Figure S11 clearly show that the titration data for L-Ala follows more steep inflection than the case of D-Ala, indicating that the formation of L-Ala adduct is favored over that of D-Ala. Also, the inflection points occur at 1.0 equivalent molar ratio in both cases, which data are also corroborate with the 1:1 binding modes in both cases.



**Figure S14:** Isothermal titration calorimetry data: (a) receptor (*S*)-1 (0.2 mM) with Lalanine (3.0 mM) (b) receptor (*S*)-1 (0.2 mM) with D-alanine (3.0 mM); measured in CH<sub>3</sub>CN at 30 °C. Alanine is used as its tetrabutylammonium carboxylate salt.





**Figure S15.** Reversibility of alanine binding to (*S*)-1. Proton-coupled <sup>19</sup>F NMR spectra of (a) receptor (*S*)-1 [1.0 mM], (b) (*S*)-1 [1.0 mM] + D-Alanine [1.5 mM], (c) L-Alanine [1.5 mM] is added to the solution of (b) which contains (*S*)-1 [1.0 mM] and D-Alanine [1.5 mM] in CD<sub>3</sub>CN, taken immediately after the addition. Alanine is used as its tetrabutylammonium carboxylate salt. (The hydrate peak observed this time is due to trace of water presented in the D-Ala sample)



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