Electronic Supplementary Information (ESI)

Enantioenrichment in sublimed amino acid mixtures

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Experimental Section:

- 1. Sublimation protocols
- 2. Powder X-ray diffraction (XRD) acquisition data
- 3. Amino acid derivatization and chiral HPLC analysis

Sublimation protocols: DL-amino acid (racemic compound, 0.03 g) and either L- or Dconfigured amino acid (0.02 g) were ground in a mortar and placed in a capped flask and heated on a hot stirring plate, which has previously attained a constant temperature of *ca.* 430 °C. Alternatively, the same enantiomeric imbalance (40%) can be attained by mixing either L- or Damino acid (0.028 g) and the opposite enantiomer (0.012 g). The mixture was stirred for a few seconds to ensure maximum contact with the hot plate. Sublimation took place suddenly and a dense cloud of condensing microcrystals could be observed for approximately 2 min. Amino acid crystals were deposited on the inner walls and covered most of the flask with the sole exception of 1.5-2 cm above the hotter base (due to the high temperature caused by conductive heating). At this stage, amino acid crystals underwent a further, slow sublimation (which took *ca.* 10 min), thus ascending the sublimation line up to 1/4 height. This process occurred without any further condensation of material from the inner atmosphere; in other words, the sublimed material was deposited on pre-existing crystals.

The solid was cooled to room temperature and then dissolved in 0.01 M HCl; the corresponding enantiomeric excesses (*ees*, referred to the whole sample on average) were measured by HPLC (*vide infra*). A gradient-like distribution of *ees* could also be inferred from analyses conducted on horizontal bands from the sublimation line upwards. The highest *ee* was measured at the sublimation line and decreased gradually as one moved upwards.

Powder XRD acquisition data: Room-temperature powder XRD patterns were collected in air using a Siemens D500 XRD diffractometer (Cu K α radiation) and a graphite monochromator. The samples were random-powder specimens. Powders were scanned over the range 2° < 2 θ < 65° at a scan rate of 0.02°/3 sec. XR Diffractograms enable phase identification of solids as racemic compounds or conglomerates. Besides valine, isoleucine converted from a racemic compound to a conglomerate phase after high-temperature sublimation (Figure S1).



Fig. S1 From top: Powder X-ray diffractograms of *rac*-isoleucine before sublimation (a racemic compound); after sublimation at 430 °C: *rac*-conglomerate (i.e. racemic mixture of individual D- and L-Ileu crystals); enantiomerically pure Ile sample for comparative purposes (bottom).

Derivatization and HPLC analysis: Enantiomeric excesses were measured via HPLC analysis with previous derivatization of amino acids. The derivatization protocol was carried out before injection and involves the treatment of the sample (2 μ L solution) with the pre-column derivatization reagent (2.2 μ L), which consists of a chiral thiol (isobutyryl-L-cysteine, 260 m*M*) and *o*-phthaldialdehyde (170 m*M*), dissolved in 1.0 *M* potassium borate buffer solution at pH 10.4.

Samples were rehydrated using 0.01 *M* HCl (750 μ L/mg) and then injected into an Agilent HPLC-1100 apparatus, equipped with a fluorescence detector. Excitation and emission wavelengths were 335 and 445 nm, respectively. A Hypersil BDS C18 reverse-phase column (5 μ m; 250 x 4 mm i.d.) was used for the analysis.

Eluent A consisted of 23 m*M* sodium acetate plus 1.5 m*M* sodium azide and 1.3 m*M* EDTA, and adjusted to pH 6.0 with 10 *M* sodium hydroxide and 10% aqueous solution of acetic acid. Eluent B was HPLC-grade methanol, while eluent C consisted of HPLC-grade acetonitrile. A linear gradient was performed at a rate of 1.0 mL min⁻¹ at 25 °C, from 95% eluent A and 5% eluent B upon injection to 76.6% eluent A, 23% eluent B, and 0.4% eluent C until min 31 and then with a progressive gradient at 1.07 mL min⁻¹ to 46.2% eluent A, 48.8% eluent B, and 5.0% eluent C until min 95.

Retention times: L-Ala (45 min), D-Ala (49 min); L-Val (60 min), D-Val (68 min); L-Ileu (72 min), D-Ileu (80 min); L-Leu (79 min), D-Leu (83 min).