## Direct observation of $\alpha$ - to $\beta$ -glycine transformation during ionic

## liquid mediated crystallization process

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## **Supplementary Information**



Figure S1 In situ powder XRD monitoring of glycine crystallization from [Bmim]PF<sub>6</sub> and [Emim]BF<sub>4</sub>. (A) 80  $\mu$ L of 50 mg/mL Aladdin glycine water solution spotted onto hydrophobic ionic liquid [Bmim]PF<sub>6</sub>. (B) 80  $\mu$ L of 50 mg/mL Sigma glycine water solution spotted onto hydrophilic ionic liquid [Emim]BF<sub>4</sub>. XRD scan time: 190 s for the each interval; 2 Theta: 5-80°.The 1' scan represents the XRD patterns of ionic liquid.

Table S1 Phase identification	rocults from th	o difforent scan	timos of a	the start of the s	estallization	of Eig	
Table ST Phase Identification	results from th	le unierent scan	times of §	giycine ch	ystamzation	ULL LIB	ure IA

Scan	Peak pos	ition (2θ)							Phases
1-4	No peaks								
5	18.904	29.764							α
6	14.729	18.930	29.734	44.185					α
7	14.717	18.938	20.085	29.745					α
8	14.768	18.934	29.781	38.351					α
9	14.749	17.869	18.941	29.783	36.221	36.603			α+ <mark>β</mark>
10	14.750	17.869	18.941	29.778	36.210	36.583	39.033		α+ <mark>β</mark>
11	14.748	17.869	18.944	23.858	29.768	36.214	36.530	39.053	α+ <mark>β</mark>
12	14.748	17.869	18.944	23.858	29.768	36.214	36.530	39.053	α+ <mark>β</mark>
13	14.748	17.869	18.944	23.858	29.768	36.214	36.530	39.053	α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively.

Table S2 Phase identification results from the different scan times of glycine crystallization of Figure 1B.

Scan	Peak pos	ition (2θ)									Phases
1-15	No peaks										
16	17.952	20.315	29.306	30.938	35.334	36.274	39.025	53.121	70.531		α+ <mark>β</mark>
17	17.892	18.964	30.113	30.982	35.316	36.203	36.559	39.026	53.005	70.588	α+ <mark>β</mark> +γ
18	17.895	18.954	30.170	30.966	35.330	36.199	36.563	39.021	53.116		α+ <mark>β</mark> +γ
19	17.889	18.935	29.253	30.162	30.981	35.324	36.192	36.636	39.026	53.117	α+ <mark>β</mark> +γ
20	17.889	18.940	29.215	30.154	30.975	35.318	36.192	36.549	39.032	53.113	α+ <mark>β</mark> +γ
21	17.890	18.967	21.837	29.242	30.157	30.974	35.337	36.195	36.632	39.026	α+ <mark>β</mark> +γ
										53.139	

The values of the peak positions marked in red, blue and black belonged to  $\beta,\gamma$  and  $\alpha$  -glycine, respectively.

Table S3 Phase identification results from the different scan times of glycine crystallization of Figure 1C.

Scan	Peak pos	sition (20)									Phases
1-5	No peaks										
6	29.908	36.360	36.452	45.440							α
7	14.897	19.118	29.888	30.982	45.438	56.749	61.908				α
8	14.910	19.115	28.507	29.906	36.280	38.583	45.439	56.733	61.915		α
9	14.909	18.005	19.120	28.497	29.906	36.308	38.585	45.434	56.733	61.916	α+ <mark>β</mark>
10	14.909	18.035	19.107	29.906	36.320	38.591	45.455	56.732	61.912		α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. The purity of  $\alpha$ -glycine was approximately 97.55% in the final products.

Table S4 Phase identification results from the different scan times of glycine crystallization of Figure 2A.

Scan	Peak posi	tion (2θ)				Phases
1'-1	No peaks					
2	61.982					α
3	14.949	29.925				α
4	17.993	29.932	33.738	36.328	39.131	α+ <mark>β</mark>
5	18.011	19.012	29.873	33.670	36.330	α+ <mark>β</mark>
6	17.987	19.024	25.326	29.850	36.333	α+ <mark>β</mark>
7	17.986	19.019	29.854	33.652	36.330	α+ <mark>β</mark>
8	17.981	19.025	29.830	33.651	36.330	α+ <mark>β</mark>
9	17.982	19.018	29.848	33.651	36.327	α+ <mark>β</mark>
10	17.981	19.029	29.837	33.651	36.329	α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. The "?" represents uncertainty of the peak.

Scan	P	eak position (	20)					Phases					
1'-2	No peaks												
3	18.013	36.357						α+ <mark>β</mark>					
4	18.019	36.321	53.083					α+ <mark>β</mark>					
5	17.989	25.329	36.303	57.794	70.632			α+ <mark>β</mark>					
6	17.978	36.303	70.609					α+ <mark>β</mark>					
7	17.976	35.402	36.302	57.748	70.585			α+ <mark>β</mark>					
8	17.975	18.986	35.455	36.303	57.773	70.624		α+ <mark>β</mark>					
9	17.975	19.076	35.440	36.302	57.762	70.635		α+ <mark>β</mark>					
10	17.976	19.069	25.328	35.380	36.302	57.856	70.720	α+ <mark>β</mark>					

Table S5 Phase identification results from the different scan times of glycine crystallization of Figure 2B.

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. It must be noticed that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the table.

Table S6 Phase identification results from the different scan times of glycine crystallization of Figure 2C.

Scan	Р	eak position (	20)				Phases
1'-2	No peaks						
3	55.866	56.559	59.345	60.875	69.240		α+ <mark>β</mark>
4	18.044	36.331	55.951				α+ <mark>β</mark>
5	18.024	36.315	39.204	55.939	76.941		α+ <mark>β</mark>
6	18.005	36.311	39.200	55.938	76.938		α+ <mark>β</mark>
7	18.003	36.296	39.190	55.694	56.356	76.938	α+ <mark>β</mark>
8	18.002	36.296	39.199	55.768	56.367	76.937	α+ <mark>β</mark>
9	18.001	36.295	39.192	55.681	56.372	76.937	α+ <mark>β</mark>
10	18.002	36.296	39.197	55.691	56.372	76.937	α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. It must be noticed that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the table.

Table S7 Phase identification results from the different scan times of glycine crystallization of Figure 2D.

Scan	Peak positi	ion (20)							Phases
1'-3	No peaks								
4	18.008	36.717							β
5	18.019	23.798	31.105	35.410	36.366	36.686	39.173	57.714	α+ <mark>β</mark>
	19.070	28.508	33.793						
6	18.007	23.788	31.094	35.392	36.340	36.675	39.173	57.666	α+ <mark>β</mark>
	19.076	28.504	33.788						
7	18.010	23.790	29.897	33.759	35.418	36.678	53.139	57.706	α+ <mark>β</mark>
	19.055	28.492	31.025	34.434	36.346	39.191			
8	18.012	23.783	29.890	33.779	35.410	36.671	56.435	57.712	α+ <mark>β</mark>
	19.081	28.498	31.085	34.407	36.327	39.153			
9	18.003	23.797	29.895	33.774	36.351	39.181	53.181	57.711	α+ <mark>β</mark>
	19.035	28.497	31.085	35.383	36.672				
10	18.010	23.794	29.883	33.795	35.405	36.664	53.181	57.749	α+ <mark>β</mark>
	19.040	28.497	31.063	34.442	36.336	39.156			

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. The numerical values marked in green indicated that the mixture phase of  $\beta$  and  $\alpha$ -glycine may be for the peak positions. It must be noticed that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the table.

Table S8 Phase identification results fr	om the different scan times of Ala	addin glycine crystallization of Figure 3A
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Scan	Peak pos	ition (2θ)							Phases		
1'-7	No peaks										
8	17.921	36.219	76.840							α+ <mark>β</mark>	
9	17.902	18.968	29.288	30.132	33.592	35.299	36.277	39.023	76.881	α+ <mark>β+γ</mark>	
10	17.903	18.977	29.282	30.156	33.606	35.292	36.227	39.026	76.885	α+ <mark>β+γ</mark>	
11	17.901	18.969	29.280	30.138	33.595	35.292	36.227	39.023	76.888	α+ <mark>β+γ</mark>	
12	17.901	18.970	29.286	30.148	33.609	35.292	36.228	39.025	76.840	α+ <mark>β+γ</mark>	
13	17.902	18.967	29.286	30.149	33.589	35.298	36.228	39.026	76.890	α+ <mark>β+γ</mark>	

The values of the peak positions marked in red, blue and black belonged to  $\beta$ ,  $\gamma$  and  $\alpha$ -glycine, respectively. It must be noticed

that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the table.

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Scan	Peak pos	ition (2θ)							Phases
1'-7	No peaks								
8	17.956	18.939	30.116	31.070	33.565	35.305	36.265	64.615	α+ <mark>β</mark> +γ
9	17.931	23.903	29.242	30.176	33.561	36.214	57.650		α+ <mark>β</mark> +γ
	18.951	28.408	29.751	31.058	35.339	56.345	64.623		
10	17.924	23.868	30.162	33.564	36.213	57.629	76.878		α+ <mark>β</mark> +γ
	18.952	29.221	31.059	35.337	56.385	64.621			
11	17.926	23.870	30.165	33.564	36.214	57.800	76.877		α+ <mark>β</mark> +γ
	18.958	29.223	31.059	35.338	56.335	64.622			
12	17.929	23.892	30.168	33.565	36.213	57.651	76.871		α+ <mark>β</mark> +γ
	18.954	29.227	31.059	35.336	56.352	64.623			

Table S9 Phase identification results from the different scan times of Sinopharm glycine crystallization of Figure 3B.

The values of the peak positions marked in red, blue and black belonged to  $\beta$ ,  $\gamma$  and  $\alpha$ -glycine, respectively.

Table S10 Phase identification results from the different scan times of Sigma glycine crystallization of Figure 3C.

Scan	Peak position (20)									
1'-6	No peaks									
7	17.933	36.201	58.849					α+ <mark>β</mark>		
8	17.946	18.929	33.639	35.310	36.270	37.949	39.032	α+ <mark>β</mark>		
9	17.958	18.960	33.629	35.284	36.295	37.963	39.027	α+ <mark>β</mark>		
10	17.972	28.439	33.630	36.293	39.035			α+ <mark>β</mark>		
	18.959	29.280	35.354	37.946						
11	17.972	28.393	33.625	36.295	39.038			α+ <mark>β</mark>		
	18.963	29.282	35.286	37.948						
12	17.975	28.381	33.627	36.294	39.039			α+ <mark>β</mark>		
	18.956	29.283	35.355	37.934						

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively.





**Figure S2** In situ monitoring of glycine crystallization from aqueous solution using powder XRD with different concentration. Individual concentrations are marked in each figure: 5.0 mg/mL s for (A-1 and -2); 6.0 mg/mL for (B-1 and -2); 10 mg/mL for (C-1 and -2); 20 mg/mL for (D-1 and -2); 50 mg/mL for (E-1 and -2). XRD program: Start angle, 5° (2 theta); End angle, 80° (2 theta); Step size, 0.0204479° (2 theta); Scan time, 381 s. 200 μL of 20.0 mg/mL Aladdin glycine solution was spotted in the sample holder for each experiment.

Table S11 Phase identification results from the different scan times of glycine solution crystallization of Figure S2A.

Scan	Peak posi	Peak position (20)								
1-9	No peaks									
10	17.826	36.126	38.983					α+ <mark>β</mark>		
11	17.819	18.850	36.156	36.547	38.998			α+ <mark>β</mark>		
12	17.821	18.813	36.154	36.517	39.006			α+ <mark>β</mark>		
13	17.821	18.864	36.177	36.527	38.965			α+ <mark>β</mark>		
14	17.818	18.805	19.978	36.135	36.496	39.004		α+ <mark>β</mark>		
15	17.818	18.791	19.964	28.335	36.162	36.537	38.947	α+ <mark>β</mark>		

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively. It must be noticed that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the tables S2A-E. It must be noticed that the values of peak position at 18.8 and 19.9°(2 $\theta$ ) belonged to  $\beta$ -glycine. The identification of peak positions is based on the two rules. One is the degree of closeness to the standard card. The other is the condition of other peaks appear in the same time.

Table S12 Phase identification results from the different scan times of glycine solution crystallization of Figure S2B.

Scan	Peak posi	tion (2θ)						Phases
1-10	No peaks							
11	17.776							β
12	17.816	18.882	29.104	36.144	36.514			α+ <mark>β</mark>
13	14.688	17.817	18.865	29.126	36.136	36.504		α+ <mark>β</mark>
14	14.690	17.817	18.880	29.122	36.165	36.524		<b>α+</b> β
15	14.680	17.718	18.872	23.618	29.123	36.147	36.577	α+ <mark>β</mark>
16	14.685	17.817	18.880	29.116	36.140	36.517		α+ <mark>β</mark>

Table S13 Phase identification results from the different scan times of glycine solution crystallization of Figure S2C.

Scan	Peak posi	ition (2θ)						Phases
1-9	No peaks							
10	14.697							α
11	17.720	18.910						β
12	14.702	17.815	18.910	20.200				α+ <mark>β</mark>
13	17.822	18.902	20.190	23.609	36.180	36.514	38.906	α+ <mark>β</mark>
14	17.820	18.901	20.198	23.618	36.208	36.486	38.908	α+ <mark>β</mark>
15	17.822	18.897	20.200	23.625	36.216	36.511	38.905	α+ <mark>β</mark>

Table S14 Phase identification results from the different scan times of glycine solution crystallization of Figure S2D.

Scan	Peak posi	tion (2θ)							Phases
1-8	No peaks								
9	17.839								β
10	17.818	20.221							β
11	17.792								β
12	14.733	17.818	18.801	20.145	33.507	35.208	36.154	36.519	α+ <mark>β</mark>
13	14.735	17.811	20.148	23.558	35.249	36.165	36.502		α+ <mark>β</mark>
14	14.727	17.812	23.594	33.546	36.159	36.507			α+ <mark>β</mark>
15	14.734	17.812	20.123	23.576	28.228	33.531	36.162	36.508	α+ <mark>β</mark>

Table S15 Phase identification results from the different scan times of glycine solution crystallization of Figure S2E.

Scan	Peak posi	ition (2θ)						Phases
1-6								
7	17.746	30.848						β
8	17.804	30.835						β
9	17.821	18.844	20.063					β
10	17.824	18.891	20.064	28.297	29.710	36.140		α+ <mark>β</mark>
11	17.814	18.873	20.117	28.301	29.652	36.148		α+ <mark>β</mark>
12	17.814	18.857	20.292	28.291	29.652	36.148		α+ <mark>β</mark>
13	17.814	18.871	20.073	28.306	29.653	36.159		α+ <mark>β</mark>
14	17.814	18.868	20.077	20.305	28.310	29.655	36.148	α+ <mark>β</mark>
15	14.674	17.813	18.879	20.064	28.314	29.668	36.148	α+ <mark>β</mark>
16	14.667	18.861	28.303	33.477	36.153	38.990	52.960	α+ <mark>β</mark>
	17.818	20.064	29.650		36.504	43.774	57.614	





Figure S3 In situ monitoring of glycine crystallization from aqueous solution using powder XRD with different volumes. Spotted volumes are marked in each figure: 1.0 µL s for (A); 5.0 µL for (B); 10 µL for (C) ; 20 µL for (D) ; 50 µL s for (E), 100 µL for (F), 200 µL for (G), 300 µL for (H). XRD program: Start angle, 5° (2 theta); End angle, 80° (2 theta); Step size, 0.0204479° (2 theta); Scan time, 381 s. 20.0 mg/mL Sigma glycine solution was used for each experiment. 26 ° and humidity 32%

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Scan	Peak position	(20)	Phases				
1-3	No peaks						
4	17.853	18.893	30.094	35.266	54.433	62.595	$\alpha + \beta + \gamma$
5	17.832	18.933	30.108	35.260	36.524	54.418	$\alpha + \beta + \gamma$
6	17.853	18.895	30.107	35.258	54.535		$\alpha + \beta + \gamma$
7	17.851	18.909	30.082	35.249	54.481		α+ <mark>β</mark> +γ

Table S16 Phase identification results from the different scan times of glycine solution crystallization of Figure S3B.

The values of the peak positions marked in red, blue and black belonged to  $\beta$ ,  $\gamma$  and  $\alpha$ -glycine, respectively. It must be noticed that the peaks were not easily found in the figures with their intensities less than 2%, which was still listed in the tables S3B-H.

Scan	Peak positio	Peak position (20)							
1-4	No peaks								
5	17.836	18.933	53.095	β					
6	17.832	18.937	53.006	β					
7	17.842	18.931	53.017	β					
8	17.843	18.933	53.006	β					
9	17.835	18.940	53.088	β					

Table S17 Phase identification results from the different scan times of glycine solution crystallization of Figure S3C.

The values of the peak positions marked in red belonged to  $\beta$ -glycine. The numerical values marked in green indicated that the phase of  $\beta$  or  $\alpha$ -glycine may be for the peak positions.

Table S18 Phase identification results from the different scan times of glycine solution crystallization of Figure S3D.

Scan	Peak positio	²eak position (2θ)								
1-5	No peaks									
6	17.860	36.210	78.780?			α+ <mark>β</mark>				
7	17.859	23.658	36.174	14.022	78.800?	α+ <mark>β</mark>				
8	17.860	23.643	36.180	78.790?		α+ <mark>β</mark>				

Table S19 Phase identification results from the different scan times of glycine solution crystallization of Figure S3E.

Scan	Peak positi	on (2 <del>0</del> )						Phases
1-2	No peaks							
3	17.779	18.858	57.562					α+ <mark>β</mark>
4	17.815	18.859	19.994	36.177	36.536			α+ <mark>β</mark>
5	14.737	17.807	18.841	23.587	36.198	36.489		α+ <mark>β</mark>
6	14.711	17.807	18.855	23.597	28.332	36.153	36.493	α+ <mark>β</mark>
7	14.712	18.853	28.305	33.562	36.169	36.514		α+ <mark>β</mark>
	17.807	23.579	<del>29.708</del>					
8	14.707	17.807	18.857	23.594	36.183			α+ <mark>β</mark>
9	14.712	18.859	28.325	29.686	33.533	38.984	36.519	α+ <mark>β</mark>
	17.810	23.619	29.089	30.892	36.178			

Table S20 Phase identification results from the different scan times of glycine solution crystallization of Figure S3F.

Scan	Peak position	(20)		Phases			
1-3	No peaks						
4	17.845	23.675	33.533	36.148	α+ <mark>β</mark>		
5	17.827	23.836	33.501	36.151	α+ <mark>β</mark>		
6	17.816	23.851	33.502	36.149	α+ <mark>β</mark>		
7	17.812	23.851	33.502	36.147	α+ <mark>β</mark>		
8	17.841	23.851	33.502	36.150	α+ <mark>β</mark>		
9	17.831	23.851	33.499	36.148	α+ <mark>β</mark>		

Scan	Peak position	Peak position (2θ)								
1-8	No peaks									
9	17.846	30.947	33.569	36.156	36.498			<b>α+</b> β		
10	17.825	28.276	30.946	33.519	36.183	36.494		α+ <mark>β</mark>		
11	17.825	28.283	30.927	33.498	36.153	36.478		α+ <mark>β</mark>		
12	17.826	28.298	30.986	33.524	36.154	36.496		<b>α+</b> β		
13	17.826	18.901	28.288	30.942	33.525	36.153	36.502	α+ <mark>β</mark>		
14	17.826	18.904	28.295	30.956	33.558	36.153		α+ <mark>β</mark>		

Table S21 Phase identification results from the different scan times of glycine solution crystallization of Figure S3G.

Table S22 Phase identification results from the different scan times of glycine solution crystallization of Figure S3H.

Scan	Peak position (20	Ð)							Phases
1-12	No peaks								
13	14.772	17.823	18.911	23.596	33.540	36.173	36.503		α+ <mark>β</mark>
14	14.694	17.819	18.892	23.589	23.809	33.521	36.180	36.578	α+ <mark>β</mark>
15	14.687	17.820	18.889	23.608	23.788	33.517	36.175	36.489	α+ <mark>β</mark>
16	14.688	17.819	18.884	23.608	23.788	33.505	36.158	36.486	α+ <mark>β</mark>





XRD program: Start angle, 5° (2 theta); End angle, 80° (2 theta); Step size, 0.0204479° (2 theta); Scan time, 762 s. No marked peaks belonged to  $\alpha$ -forms glycine. Phase identifications of glycine crystals were compared with standard cards of PDF#06-0230 ( $\gamma$ -glycine), PDF#32-1702 ( $\alpha$ -glycine) and PDF#02-0171 ( $\beta$ -glycine).

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Brands	Peak positions(2θ)					Phases			
Sigma	14.832	18.975	20.134	23.913	28.407	29.848	35.421		α
	36.437	38.680	40.859	45.393	29.220				
Aladdin	14.842	19.072	20.190	23.895	28.503	29.351	29.810	30.220	α+γ
	31.107	35.257	35.264	36.210	36.637	37.884	38.384	38.721	
	40.464	40.847	42.101	42.926	45.355	47.334	52.475	53.910	
	55.372	56.452	58.886	61.035	77.382	α+β			
Sinopharm	14.932	19.173	20.289	24.068	28.541	29.367	29.800	30.284	α+γ
	31.192	35.442	36.062	36.540	38.000	38.587	40.793	41.035	
	42.251	43.036	44.229	47.012	47.388	50.552	56.048	56.319	
	62.710	69.528	74.781	75.191	77.372				

Table S23 Phase identification results from XRD patterns of three commercial brands of Figure S4.

The values of the peak positions marked in blue and black belonged to  $\gamma$  and  $\alpha$ -glycine, respectively.





Figure S5 In situ monitoring of glycine crystallization from aqueous solution using powder XRD with three commercial brands. (A) Aladdin; (B)Sinopharm;(C) Sigma. XRD program: Start angle, 5° (2 theta); End angle, 80°(2 theta); Step size, 0.02° (2 theta); Scan time, 381 s. 400  $\mu$ L of 20.0 mg/mL glycine solution was used for each experiment at 26 ° and humidity 32%.

Table S24 Phase identification results from the different scan times of Figure S5A.

Scan		Peak position (20)				
1-18	No peaks					
19	17.845	33.545	36.169	36.518		α+ <mark>β</mark>
20	17.828	18.868	33.518	36.155	36.509	α+ <mark>β</mark>
21	17.828	18.868	33.519	36.154	36.501	α+ <mark>β</mark>
22	17.828	18.871	33.526	36.155	36.509	α+ <mark>β</mark>
23	17.829	18.842	33.530	36.156	36.511	α+ <mark>β</mark>
24	17.829	18.871	33.534	36.154	36.508	α+ <mark>β</mark>
25	17.828	18.872	33.497	36.154	36.500	α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$  -glycine, respectively.

Table S25 Phase identification results from the different scan times of Figure S5B.

Scan	Peak position (2θ)						Phases
1-18	No peaks						
19	14.816						α
20	14.785	17.838	33.519	36.166			α+ <mark>β</mark>
21	14.758	17.842	18.911	28.302	33.503	36.167	α+ <mark>β</mark>
22	14.741	17.841	18.899	28.351	33.516	36.167	α+ <mark>β</mark>
23	14.752	17.838	18.897	28.339	33.511	36.168	α+ <mark>β</mark>
24	14.752	17.838	18.902	28.329	33.516	36.168	α+ <mark>β</mark>
25	14.750	17.839	18.906	28.324	33.515	36.171	α+ <mark>β</mark>
26	14.744	17.838	18.900	28.346	33.501	36.170	α+ <mark>β</mark>

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha$ -glycine, respectively.

Scan	Peak posi	Peak position (20)				
1-14	No peaks					
15	28.345	29.704	41.565			α
16	17.778	23.639	29.632	33.535	76.828	α+ <mark>β</mark>
	18.918	28.346	30.913	36.148		
17	14.717	18.923	28.300	33.535	49.000	α+ <mark>β</mark>
	17.848	23.832	29.738	36.166	55.354	
18	14.718	18.923	28.299	33.535	49.001	α+ <mark>β</mark>
	17.849	23.834	29.731	36.157	55.348	
19	14.718	18.924	28.299	33.536	48.996	α+ <mark>β</mark>
	17.849	23.832	29.729	36.166	55.348	
20	14.719	18.910	28.299	33.535	48.998	α+ <mark>β</mark>
	17.849	23.834	29.731	36.157	55.347	
21	14.715	18.910	28.299	33.535	48.997	α+ <mark>β</mark>
	17.849	23.833	29.731	36.166	55.346	
22	14.719	18.910	28.299	33.535	48.999	α+ <mark>β</mark>
	17.848	23.832	29.731	36.157	55.348	
23	14.718	18.911	28.299	33.535	49.001	α+ <mark>β</mark>
	17.849	23.834	29.730	36.166	55.348	

Table S26 Phase identification results from the different scan times of Figure S5C.

The values of the peak positions marked in red and black belonged to  $\beta$  and  $\alpha\mbox{-glycine},$  respectively.

Concentration	Volume	Ambient	Substrate	Crystallization	Solution	Ref.
		conditions		Time		
4.0-35.9 g/mL	unknown	25°C	Crystallizing dishes	unknown	Methanol or	42
					Ethanol-	
					water	
10 g/30 mL	unknown	room	Crystallizing dishes	unknown	α-amino	43
		temperature			acids in	
					water	
Supersaturated	unknown	25°C	Macroemulsions	5 h	water in oil	44
30.78 g/100 g	unknown	80°C	Emulsions	unknown	water in oil	45
3.0 M	unknown	40 °C to 2°C	SAMs/Au/Si	12.6-633.0h	Pure water	46
				or >200		
3.2 M	unknown	23°C, 35%	SAMs/Au/Si	5-450 min	Pure water	47
1-133 mg/mL	50 μL	21°C, 30%	Pt/SiO2/Si	unknown	Pure water	48
18%	unknown	vacuum	p-PS-	6 h	Pure water	49
			PDMA monoliths			
5% -18%	unknown	vacuum	CPG and p-PS-	6 h	Pure water	50
			PDMA monoliths			
10-17.5%	unknown	90-180°C	unknown	unknown	Pure water	51
16.6 mg/mL	unknown	dry air	unknown	unknown	Pure water	52
5-50 mg/mL	1-400 μL	26°C, 32%	XRD glass sample	30-150 min	Pure water	This
(0.0666-0.666M)			holder		or ILs-water	paper

## Table S27 Crystallization condition of $\beta$ -glycine from references 42-52

Brand	Solvent	The percentage	Figure
		of β-glycine	
Aladdin	Water	45.71%	Figure 1A
Aladdin	[Bmim]PF <sub>6</sub> -water	35.50%	Figure 1B
Aladdin	[Emim]BF <sub>4</sub> - water	86.38%	Figure 2A
Aladdin	[Emim]BF <sub>4</sub> - water	89.89%	Figure 2B
Aladdin	[Emim]BF <sub>4</sub> - water	60.34%	Figure 2C
Aladdin	[Emim]BF <sub>4</sub> - water	56.42%	Figure 2D
Aladdin	[Bmim]PF <sub>6</sub> -water	41.33%	Figure 3A
Sigma	[Emim]BF <sub>4</sub> - water	2.45%	Figure 1C
Sigma	[Bmim]PF <sub>6</sub> -water	55.96%	Figure 3C
Sigma	Water	62.46%	Figure S5C

Table S28 The percentage of  $\beta\mbox{-glycine}$  crystals from different crystallization condition

The percentage of  $\beta$ -glycine was calculated according to the peak areas.