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Structural data for L-leucine hydrochloride monohydrate:

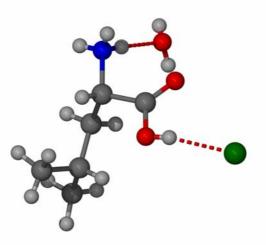
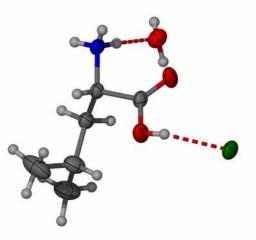


Figure 1. Ball and stick representation of the asymmetric unit found in the crystal structure of  $_{L}$ -leucine hydrochloride monohydrate showing hydrogen bonding between the structural components.



**Figure 2.** The asymmetric unit found in the crystal structure of  $_{L}$ -leucine hydrochloride monohydrate, anisotropic displacement parameters shown at the 50% level. Hydrogen bonding between structural components is also shown.

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**Table 1.** Crystal data and structure refinement for  $_L$ -leucine hydrochloride monohydrate.

Identification code	L-Leucine hydrochloride monohydrate	
Formula	C <sub>6</sub> H <sub>16</sub> CINO <sub>3</sub>	
Formula weight	185.65	
Size	0.46 x 0.18 x 0.08 mm	
Crystal morphology	Colourless plate	
Temperature	150(2) K	
Wavelength	0.71073 Å [Mo- <i>K</i> <sub>α</sub> ]	
Crystal system	Orthorhombic	
Space group	P212121	
Unit cell dimensions	$a = 6.2563(13)$ Å $\alpha = 90^{\circ}$	
	$b = 6.3951(13) \text{ Å} \qquad \beta = 90^{\circ}$	
	$c = 26.865(5) \text{ Å} \qquad \gamma = 90^{\circ}$	
Volume	1074.9(4) Å <sup>3</sup>	
Ζ	4	
Density (calculated)	1.147 Mg/m <sup>3</sup>	
Absorption coefficient	0.325 mm <sup>-1</sup>	
<i>F</i> (000)	400	
Data collection range	$3.34 \le \theta \le 27.5^{\circ}$	
Index ranges	$-8 \le h \le 4$ , $-7 \le k \le 4$ , $-11 \le l \le 33$	
Reflections collected	1746	
Independent reflections	1473 [ $R(int) = 0.0678$ ]	
Observed reflections	1299 [ <i>I</i> >2σ( <i>I</i> )]	
Absorption correction	multi-scan	
Max. and min. transmission	0.949 and 0.104	
Refinement method	Full	
Data / restraints / parameters	1473 / 0 / 164	
Goodness of fit	1.033	
Final <i>R</i> indices $[I \ge 2\sigma(I)]$	$R_1 = 0.0343, wR_2 = 0.0777$	
<i>R</i> indices (all data)	$R_1 = 0.0418, wR_2 = 0.0818$	

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Largest diff. peak and hole	0.159 and -0.167e.Å <sup>-3</sup>
Absolute structure parameter	-0.01(9)