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## **Potassium *N*-(6-oxocyclohexa-1,3-dien-5-ylidenemethyl)glycinate**

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### **Electronic paper**

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Potassium *N*-(6-oxocyclohexa-1,3-dien-5-ylidene)methyl)glycinateVerónica Paredes-García,<sup>a\*</sup> Diego Venegas-Yazigi,<sup>a</sup>  
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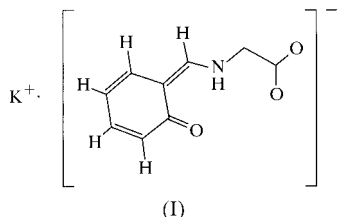
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We are very interested in characterizing ligands derived from salicylaldehyde and  $\alpha$ -amino acids. These ligands form interesting complexes with different transition metals, which are very useful for understanding some biological systems. For



Schiff bases derived from salicylaldehyde and different alkylamines or arylamines, the enol-imine form is predominant. The title compound, (I), shows bond distances which are clearly in agreement with the keto-amine form.

## Experimental

## Crystal data

 $K^+ \cdot C_9H_8NO_3^-$  $M_r = 217.26$ Orthorhombic, *Pcca* $a = 13.8864$  (2) Å $b = 18.4943$  (7) Å $c = 6.9664$  (9) Å $V = 1789.1$  (2) Å<sup>3</sup> $Z = 8$  $D_x = 1.613$  Mg m<sup>-3</sup>Mo  $K\alpha$  radiationCell parameters from 18191  
reflections $\theta = 2.93$ – $27.49^\circ$  $\mu = 0.570$  mm<sup>-1</sup> $T = 100$  (1) K

Plate, orange

 $0.30 \times 0.30 \times 0.10$  mm

## Data collection

KappaCCD diffractometer

 $\varphi$  and  $\omega$  scans with  $\kappa$  offsetsAbsorption correction: multi-scan  
(*DENZO-SMN*; Otwinowski &  
Minor, 1997) $T_{\min} = 0.848$ ,  $T_{\max} = 0.945$ 

18 191 measured reflections

2036 independent reflections

1580 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.065$  $\theta_{\max} = 27.49^\circ$  $h = -18 \rightarrow 18$  $k = -24 \rightarrow 24$  $l = -8 \rightarrow 8$ 

Intensity decay: none

## Refinement

Refinement on  $F^2$  $R(F) = 0.034$  $wR(F^2) = 0.091$  $S = 1.051$ 

2036 reflections

131 parameters

H atoms treated by a mixture of  
independent and constrained  
refinement $w = 1/[\sigma^2(F_o^2) + (0.0531P)^2$   
 $+ 0.0850P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} = 0.001$  $\Delta\rho_{\max} = 0.312$  e Å<sup>-3</sup> $\Delta\rho_{\min} = -0.44$  e Å<sup>-3</sup>

Table 1

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1N...O3	0.91 (2)	1.91 (2)	2.6310 (19)	135.4 (18)

Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXTL/PC* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL/PC*; software used to prepare material for publication: *SHELXTL/PC*.

Data were collected at the University of Toronto (where all structural calculations were carried out) on a Nonius KappaCCD purchased with funds from NSERC Canada. We would like to thank FONDECYT for funding of Project 2960043.

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