

# Structural characterization of form I of anhydrous rifampicin

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## Supporting Information

**Table S1 - Final coordinates and equivalent isotropic displacement parameters ( $\mathbf{U}_{\text{iso}} = \mathbf{B}_{\text{iso}}/8\pi^2$ ) for all atoms in the polymorph I of anhydrous rifampicin crystal structure.**

Atom	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}$ ( $\text{\AA}^2$ )
O(2)	0.2744(9)	0.300(2)	1.0773(19)	0.069(1)
O(6)	0.1354(12)	0.0768(2)	0.841(2)	0.069(1)
O(9)	0.0525(11)	0.175(3)	0.791(2)	0.069(1)
O(14)	0.0264(10)	0.503(3)	0.7699(18)	0.069(1)
O(17)	0.0571(11)	0.595(3)	0.6599(19)	0.069(1)
O(21)	0.0707(11)	0.724(3)	0.590(2)	0.069(1)
O(30)	0.3302(10)	0.670(2)	0.801(2)	0.069(1)
O(34)	0.3881(9)	0.535(2)	0.9642(18)	0.069(1)
O(38)	0.3664(12)	0.289(3)	0.968(2)	0.069(1)
O(41)	0.4371(11)	0.366(2)	1.119(2)	0.069(1)
O(55)	0.1095(10)	0.271(3)	0.5205(19)	0.069(1)
O(59)	0.1020(10)	0.128(2)	0.611(2)	0.069(1)
N(19)	0.0944(13)	0.609(3)	0.507(3)	0.069(1)
N(46)	0.1348(13)	0.511(3)	0.403(3)	0.069(1)
N(47)	0.1595(13)	0.504(4)	0.340(3)	0.069(1)
N(50)	0.1512(15)	0.478(3)	0.138(3)	0.069(1)
C(1)	0.3105(11)	0.250(3)	1.174(2)	0.069(1)
C(3)	0.261(2)	0.250(3)	0.982(3)	0.069(1)
C(4)	0.2025(18)	0.198(4)	0.938(3)	0.069(1)
C(5)	0.1891(19)	0.125(3)	0.875(3)	0.069(1)
C(7)	0.078(2)	0.111(4)	0.747(4)	0.069(1)
C(8)	0.0363(11)	0.027(3)	0.693(2)	0.069(1)
C(10)	0.0513(18)	0.261(5)	0.749(6)	0.069(1)
C(11)	0.0365(17)	0.341(6)	0.784(4)	0.069(1)
C(12)	0.0133(13)	0.339(3)	0.860(3)	0.069(1)
C(13)	0.0386(19)	0.424(5)	0.735(5)	0.069(1)
C(15)	0.063(2)	0.429(6)	0.665(4)	0.069(1)
C(16)	0.0676(18)	0.517(4)	0.621(4)	0.069(1)
C(18)	0.0951(17)	0.522(4)	0.559(4)	0.069(1)
C(20)	0.0967(17)	0.699(4)	0.543(4)	0.069(1)
C(22)	0.1122(17)	0.771(4)	0.486(4)	0.069(1)
C(23)	0.0597(12)	0.834(3)	0.405(3)	0.069(1)
C(24)	0.1661(14)	0.771(3)	0.494(3)	0.069(1)
C(25)	0.2143(16)	0.702(3)	0.547(3)	0.069(1)
C(26)	0.2693(17)	0.712(3)	0.563(3)	0.069(1)
C(27)	0.3237(17)	0.648(4)	0.628(4)	0.069(1)
C(28)	0.3283(10)	0.576(3)	0.551(3)	0.069(1)
C(29)	0.3266(16)	0.599(4)	0.727(3)	0.069(1)
C(31)	0.3792(17)	0.532(4)	0.788(3)	0.069(1)
C(32)	0.4411(11)	0.582(3)	0.842(2)	0.069(1)
C(33)	0.3773(18)	0.475(4)	0.878(3)	0.069(1)
C(35)	0.3194(19)	0.416(4)	0.837(3)	0.069(1)
C(36)	0.3052(11)	0.351(3)	0.741(2)	0.069(1)
C(37)	0.3209(16)	0.362(4)	0.931(3)	0.069(1)
C(39)	0.4238(19)	0.303(3)	1.058(3)	0.069(1)
C(40)	0.4580(13)	0.212(3)	1.094(3)	0.069(1)
C(42)	0.2598(15)	0.314(3)	0.894(3)	0.069(1)
C(43)	0.2095(11)	0.388(3)	0.855(2)	0.069(1)
C(44)	0.1080(18)	0.438(5)	0.522(4)	0.069(1)
C(45)	0.1283(16)	0.436(3)	0.445(4)	0.069(1)

C(48)	0.1581(16)	0.411(3)	0.297(3)	0.069(1)
C(49)	0.1849(14)	0.411(3)	0.227(3)	0.069(1)
C(51)	0.1720(13)	0.475(3)	0.061(3)	0.069(1)
C(52)	0.1613(16)	0.571(3)	0.187(3)	0.069(1)
C(53)	0.1350(14)	0.578(4)	0.258(3)	0.069(1)
C(54)	0.1005(17)	0.351(4)	0.560(4)	0.069(1)
C(56)	0.080(2)	0.345(6)	0.634(5)	0.069(1)
C(57)	0.071(2)	0.259(4)	0.673(5)	0.069(1)
C(58)	0.0830(18)	0.167(5)	0.662(4)	0.069(1)
H(60)	0.2158(19)	0.105(3)	0.853(3)	0.082(1)
H(61)	0.0299(11)	-0.003(3)	0.746(2)	0.082(1)
H(62)	-0.0024(11)	0.046(3)	0.629(2)	0.082(1)
H(63)	0.0550(11)	-0.017(3)	0.668(2)	0.082(1)
H(64)	0.1736(18)	0.219(4)	0.956(3)	0.082(1)
H(65)	0.293(2)	0.205(3)	1.005(3)	0.082(1)
H(66)	0.2487(15)	0.276(3)	0.830(3)	0.082(1)
H(67)	0.0433(13)	0.311(3)	0.929(3)	0.082(1)
H(68)	0.0053(13)	0.402(3)	0.873(3)	0.082(1)
H(69)	-0.0245(13)	0.304(3)	0.825(3)	0.082(1)
H(70)	0.3319(16)	0.404(4)	0.991(3)	0.082(1)
H(71)	0.2174(11)	0.425(3)	0.917(2)	0.082(1)
H(72)	0.2088(11)	0.428(3)	0.799(2)	0.082(1)
H(73)	0.1704(11)	0.358(3)	0.823(2)	0.082(1)
H(74)	0.3481(11)	0.239(3)	1.178(2)	0.082(1)
H(75)	0.3197(11)	0.287(3)	1.237(2)	0.082(1)
H(76)	0.2946(11)	0.189(3)	1.176(2)	0.082(1)
H(77)	0.1045(10)	0.209(3)	0.5475(19)	0.082(1)
H(78)	0.0277(10)	0.564(3)	0.7369(18)	0.082(1)
H(79)	0.2857(19)	0.460(4)	0.808(3)	0.082(1)
H(80)	0.1369(16)	0.377(3)	0.425(4)	0.082(1)
H(81)	0.0239(11)	0.639(3)	0.6091(19)	0.082(1)
H(82)	0.4117(18)	0.432(4)	0.911(3)	0.082(1)
H(83)	0.2653(11)	0.324(3)	0.710(2)	0.082(1)
H(84)	0.3049(11)	0.388(3)	0.684(2)	0.082(1)
H(85)	0.3365(11)	0.305(3)	0.766(2)	0.082(1)
H(86)	0.0907(13)	0.604(3)	0.442(3)	0.082(1)
H(87)	0.3763(17)	0.488(4)	0.734(3)	0.082(1)
H(88)	0.3885(9)	0.510(2)	1.0293(18)	0.082(1)
H(89)	0.4603(13)	0.189(3)	1.033(3)	0.082(1)
H(90)	0.5000(13)	0.224(3)	1.154(3)	0.082(1)
H(91)	0.4379(13)	0.166(3)	1.112(3)	0.082(1)
H(92)	0.2890(16)	0.564(4)	0.697(3)	0.082(1)
H(93)	0.4740(11)	0.537(3)	0.875(2)	0.082(1)
H(94)	0.4420(11)	0.618(3)	0.786(2)	0.082(1)
H(95)	0.4460(11)	0.625(3)	0.899(2)	0.082(1)
H(96)	0.1453(14)	0.638(4)	0.293(3)	0.082(1)
H(97)	0.0909(14)	0.571(4)	0.209(3)	0.082(1)
H(98)	0.1164(16)	0.389(3)	0.251(3)	0.082(1)
H(99)	0.1822(16)	0.368(3)	0.358(3)	0.082(1)
H(100)	0.3595(17)	0.687(4)	0.660(4)	0.082(1)
H(101)	0.3322(10)	0.649(2)	0.869(2)	0.082(1)
H(102)	0.1740(14)	0.823(3)	0.461(3)	0.082(1)
H(103)	0.0448(12)	0.867(3)	0.444(3)	0.082(1)
H(104)	0.0738(12)	0.880(3)	0.375(3)	0.082(1)
H(105)	0.0266(12)	0.798(3)	0.347(3)	0.082(1)
H(106)	0.2046(16)	0.584(3)	0.231(3)	0.082(1)
H(107)	0.1421(16)	0.616(3)	0.128(3)	0.082(1)
H(108)	0.1827(14)	0.351(3)	0.197(3)	0.082(1)

H(109)	0.2274(14)	0.431(3)	0.273(3)	0.082(1)
H(110)	0.2744(17)	0.763(3)	0.526(3)	0.082(1)
H(111)	0.2947(10)	0.532(3)	0.522(3)	0.082(1)
H(112)	0.3263(10)	0.610(3)	0.493(3)	0.082(1)
H(113)	0.3674(10)	0.544(3)	0.595(3)	0.082(1)
H(114)	0.2062(16)	0.646(3)	0.574(3)	0.082(1)
H(115)	0.2154(13)	0.490(3)	0.102(3)	0.082(1)
H(116)	0.1656(13)	0.414(3)	0.030(3)	0.082(1)
H(117)	0.1494(13)	0.520(3)	0.002(3)	0.082(1)

**Table S2 Some selected H-bonds in the structure of form I of anhydrous RMP. Intermolecular bonds are denoted by superscripts.**

D–H···A	D–H (Å)	H···A (Å)	D···A (Å)	D–H···A (°)
O(55)–H(77)···O(59)	1.00(6)	1.49(5)	2.48(5)	167(3)
O(14)–H(78)···O(17)	1.00(6)	1.70(4)	2.49(5)	132(4)
O(17)–H(81)···N(19) <sup>a</sup>	0.99(5)	2.60(5)	3.31(5)	128(4)
N(19)–H(86)···N(46)	0.88(5)	2.02(6)	2.64(6)	126(4)
O(34)–H(88)···O(41)	0.99(3)	2.39(4)	3.05(4)	123(3)
O(34)–H(88)···O(6) <sup>b</sup>	0.99(3)	2.44(4)	3.21(4)	134(3)
O(30)–H(101)···O(34)	0.99(4)	2.11(4)	2.76(4)	121(3)
C(3)–H(65)···O(38)	0.96(7)	2.54(6)	2.90(7)	103(4)
C(12)–H(68)···O(14)	0.97(6)	2.33(5)	2.78(6)	108(4)
C(37)–H(70)···O(34)	0.95(6)	2.53(6)	2.91(6)	104(4)
C(37)–H(70)···O(41)	0.95(6)	2.37(5)	2.73(5)	102(5)
C(45)–H(80)···O(55)	0.96(6)	2.39(6)	2.75(6)	102(5)
C(25)–H(114)···N(19)	0.96(6)	2.56(6)	3.13(6)	118(4)

<sup>a</sup>Symmetry: -x, y, 1-z; <sup>b</sup>Symmetry: ½-x, ½+y, 2-z.

**Table S3 - Some selected bond distances for the polymorph I of anhydrous rifampicin.**

Bonds	Lengths (Å)
O(2)-C(1)	1.37(4)
O(2)-C(3)	1.40(5)
O(6)-C(5)	1.38(6)
O(6)-C(7)	1.44(6)
O(9)-C(7)	1.46(7)
O(9)-C(10)	1.36(8)
O(14)-C(13)	1.34(8)
O(17)-C(16)	1.34(7)
O(21)-C(20)	1.23(6)
O(30)-C(29)	1.43(6)
O(34)-C(33)	1.40(5)

O(38)-C(37)	1.44(6)
O(38)-C(39)	1.35(5)
O(41)-C(39)	1.17(5)
O(55)-C(54)	1.35(7)
O(59)-C(58)	1.21(7)
N(19)-C(18)	1.44(7)
N(19)-C(20)	1.38(7)
N(46)-N(47)	1.36(6)
N(46)-C(45)	1.28(6)
N(47)-C(53)	1.44(7)
N(47)-C(48)	1.46(7)
N(50)-C(51)	1.47(6)
C(33)-C(35)	1.53(7)
C(35)-C(37)	1.53(6)
C(35)-C(36)	1.53(6)
C(37)-C(42)	1.53(7)
C(39)-C(40)	1.50(6)
C(42)-C(43)	1.53(6)
C(44)-C(45)	1.45(7)
C(44)-C(54)	1.41(9)
C(48)-C(49)	1.50(6)
C(52)-C(53)	1.50(6)
C(54)-C(56)	1.42(8)
C(56)-C(57)	1.42(10)
C(57)-C(58)	1.38(9)
N(50)-C(52)	1.46(6)
N(50)-C(49)	1.45(6)
C(3)-C(4)	1.49(7)
C(3)-C(42)	1.54(6)
C(4)-C(5)	1.30(7)
C(7)-C(8)	1.52(7)
C(7)-C(58)	1.52(8)
C(10)-C(57)	1.43(9)
C(10)-C(11)	1.38(10)

C(11)-C(12)	1.50(7)
C(11)-C(13)	1.39(10)
C(13)-C(15)	1.45(8)
C(15)-C(56)	1.43(11)
C(15)-C(16)	1.44(10)
C(16)-C(18)	1.41(8)
C(18)-C(44)	1.42(9)
C(20)-C(22)	1.50(8)
C(22)-C(23)	1.51(7)
C(22)-C(24)	1.34(7)
C(24)-C(25)	1.44(6)
C(25)-C(26)	1.32(7)
C(26)-C(27)	1.50(7)
C(27)-C(29)	1.54(7)
C(27)-C(28)	1.56(7)
C(29)-C(31)	1.50(7)
C(31)-C(33)	1.55(7)
C(31)-C(32)	1.53(6)

**Table S4 - Some selected bond angles for the polymorph I of anhydrous rifampicin.**

Bonds	Angles (°)
C(1)-O(2)-C(3)	113(3)
C(5)-O(6)-C(7)	119(3)
C(7)-O(9)-C(10)	107(4)
C(37)-O(38)-C(39)	120(4)
C(18)-N(19)-C(20)	129(4)
N(47)-N(46)-C(45)	118(5)
N(46)-N(47)-C(53)	109(4)
C(48)-N(47)-C(53)	115(3)
N(46)-N(47)-C(48)	116(4)
C(49)-N(50)-C(52)	109(3)
C(51)-N(50)-C(52)	110(4)
C(49)-N(50)-C(51)	110(4)
O(2)-C(3)-C(4)	109(4)

O(2)-C(3)-C(42)	112(3)
C(4)-C(3)-C(42)	113(3)
C(3)-C(4)-C(5)	122(5)
O(6)-C(5)-C(4)	119(5)
C(8)-C(7)-C(58)	111(4)
O(6)-C(7)-O(9)	107(3)
O(6)-C(7)-C(8)	107(4)
O(6)-C(7)-C(58)	116(5)
C(25)-C(26)-C(27)	127(4)
C(26)-C(27)-C(28)	112(4)
C(26)-C(27)-C(29)	114(4)
C(28)-C(27)-C(29)	111(4)
C(27)-C(29)-C(31)	114(4)
O(30)-C(29)-C(27)	108(4)
O(30)-C(29)-C(31)	111(3)
C(29)-C(31)-C(32)	112(4)
C(29)-C(31)-C(33)	113(4)
C(32)-C(31)-C(33)	109(3)
O(34)-C(33)-C(31)	109(4)
O(34)-C(33)-C(35)	111(4)
C(31)-C(33)-C(35)	116(3)
C(33)-C(35)-C(36)	113(4)
C(36)-C(35)-C(37)	111(4)
C(33)-C(35)-C(37)	113(3)
O(38)-C(37)-C(35)	108(4)
C(35)-C(37)-C(42)	113(3)
O(38)-C(37)-C(42)	107(4)
O(38)-C(39)-C(40)	110(4)
O(41)-C(39)-C(40)	122(4)
O(38)-C(39)-O(41)	123(5)
C(3)-C(42)-C(37)	115(3)
C(3)-C(42)-C(43)	110(4)
C(37)-C(42)-C(43)	109(4)
C(18)-C(44)-C(45)	124(5)

C(18)-C(44)-C(54)	119(5)
C(45)-C(44)-C(54)	117(5)
O(9)-C(7)-C(8)	111(4)
O(9)-C(7)-C(58)	106(5)
O(9)-C(10)-C(11)	122(6)
O(9)-C(10)-C(57)	113(5)
C(11)-C(10)-C(57)	125(6)
C(10)-C(11)-C(12)	123(6)
C(10)-C(11)-C(13)	115(5)
C(12)-C(11)-C(13)	121(6)
O(14)-C(13)-C(11)	117(5)
O(14)-C(13)-C(15)	120(6)
C(11)-C(13)-C(15)	123(6)
C(13)-C(15)-C(16)	121(7)
C(13)-C(15)-C(56)	120(7)
C(16)-C(15)-C(56)	119(5)
O(17)-C(16)-C(18)	120(5)
C(15)-C(16)-C(18)	121(5)
O(17)-C(16)-C(15)	118(5)
C(16)-C(18)-C(44)	120(5)
N(19)-C(18)-C(44)	118(4)
N(19)-C(18)-C(16)	120(5)
O(21)-C(20)-N(19)	123(5)
O(21)-C(20)-C(22)	119(5)
N(19)-C(20)-C(22)	114(4)
C(20)-C(22)-C(23)	116(4)
C(20)-C(22)-C(24)	122(5)
C(23)-C(22)-C(24)	122(4)
C(22)-C(24)-C(25)	128(4)
C(24)-C(25)-C(26)	124(4)
N(46)-C(45)-C(44)	122(5)
N(47)-C(48)-C(49)	112(4)
N(50)-C(49)-C(48)	108(4)
N(50)-C(52)-C(53)	111(4)

N(47)-C(53)-C(52)	110(4)
C(44)-C(54)-C(56)	122(6)
O(55)-C(54)-C(44)	120(5)
O(55)-C(54)-C(56)	119(5)
C(15)-C(56)-C(57)	118(6)
C(15)-C(56)-C(54)	119(7)
C(54)-C(56)-C(57)	124(7)
C(10)-C(57)-C(56)	119(6)
C(10)-C(57)-C(58)	108(5)
C(56)-C(57)-C(58)	133(6)
O(59)-C(58)-C(7)	119(6)
O(59)-C(58)-C(57)	134(6)
C(7)-C(58)-C(57)	107(5)

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