

Glyoxal (グリオキサール)

Chemical Name: Glyoxal  
 Synonym: 1,2-Ethanedione  
 Ethanedial  
 Diformyl  
 Glyoxalaldehyde  
 Molecular weight: 58.04  
 Melting point: 15°C  
 Boiling point: 51°C  
 Chemical Structure  

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      CHO
    
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 CAS No : 107-22-2  
 MITI No : (2)-509, (2)-510  
 Source of Substance: Tokyo Kasei Kogyo Co. Ltd  
 Lot.No. : AY01  
 Purity: 40 % (in water)  
 Vehicle: H<sub>2</sub>O

Mutagenicity  
 in Bacterial Test: Positive

IARC Evaluation: not yet cited

Con. μg/ plate	Experimental Data									
	Number of Revertants/plate									
	Base-substitution						Frame-shift			
	TA100		TA1535		WP2uvrA		TA98		TA1537	
	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
H <sub>2</sub> O	(158)	(166)	(7)	(8)	(24)	(26)	(23)	(23)	(6)	(8)
	192	185	3	7	18	28	20	22	7	7
7.81	(185)	(197)	(6)	(8)	(20)	(28)	(23)	(21)	(7)	(7)
	177	209	8	8	22	27	25	20	6	7
	213	221	13	15	24	23	22	22	6	13
15.6	(211)	(224)	(12)	(13)	(24)	(29)	(18)	(21)	(7)	(14)
	208	227	10	10	23	34	13	20	7	15
	332	255	7	9	27	42	23	23	3	9
31.3	(315)	(268)	(8)	(11)	(25)	(41)	(20)	(23)	(5)	(9)
	298	281	8	13	22	39	16	23	6	9
	492	436	10	20	25	32	24	28	7	10
62.5	(458)	(426)	(12)	(19)	(24)	(35)	(25)	(25)	(7)	(11)
	423	415	13	18	22	37	25	21	6	12
	562	716	10	14	51	73	61	24	7	8
125	(557)	(721)	(10)	(14)	(51)	(69)	(52)	(32)	(7)	(9)
	551	726	10	13	50	65	43	39	6	9
	596*	853*	8*	12*	71	105	46*	42	16*	13
250	(546*)	(838*)	(8*)	(13*)	(67)	(102)	(41*)	(43)	(15*)	(15)
	495*	822*	8*	14*	62	99	35*	43	13*	16
	298*	325*	10*	12*	39*	106*	6*	47*	5*	13*
500	(280*)	(307*)	(10*)	(11*)	(41*)	(106*)	(4*)	(44*)	(3*)	(13*)
	262*	289*	9*	10*	43*	105*	1*	40*	1*	12*
	113*	418*	7*	16*	30*	38*	0*	9*	0*	8*
1000	(169*)	(414*)	(6*)	(16*)	(35*)	(34*)	(0*)	(12*)	(0*)	(11*)
	224*	409*	5*	15*	39*	29*	0*	15*	0*	14*
Judgement	+	+	-	+	+	+	+	-	+	-
Specific Mutagenicity	4800	4440		176	216	344	232		36.0	
Positive	AF2	2AA	NaN <sub>3</sub>	2AA	AF2	2AA 20	AF2	2AA	9AA	2AA
Control	(667)	(835)	(241)	(208)	(223)	(1111)	(373)	(413)	(437)	(149)

Experimental Data

Con. $\mu$ g/ plate	Number of Revertants/plate									
	Base-substitution						Frame-shift			
	TA100		TA1535		WP2uvrA		TA98		TA1537	
	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
H <sub>2</sub> O	( 178 )	( 186 )	( 7 )	( 8 )	( 28 )	( 29 )	( 28 )	( 28 )	( 8 )	( 10 )
	190	197	5	7	21	25	31	30	5	10
	183	193	10	10	20	32	23	20	3	9
10	( 187 )	( 195 )	( 8 )	( 9 )	( 21 )	( 29 )	( 27 )	( 25 )	( 4 )	( 10 )
	251	217	6	13	27	31	25	27	6	9
	297	228	6	12	27	32	29	30	10	12
20	( 274 )	( 223 )	( 6 )	( 13 )	( 27 )	( 32 )	( 27 )	( 29 )	( 8 )	( 11 )
	414	318	10	15	30	40	31	37	10	7
	466	329	8	16	25	34	35	25	7	9
40	( 440 )	( 324 )	( 9 )	( 16 )	( 28 )	( 37 )	( 33 )	( 31 )	( 9 )	( 8 )
	704	420	10	20	24	45	31	30	6	10
	644	497	10	16	29	44	38	32	10	12
60	( 674 )	( 459 )	( 10 )	( 18 )	( 27 )	( 45 )	( 35 )	( 31 )	( 8 )	( 11 )
	773	579	9	21	25	59	50	34	6	10
	803	570	12	16	29	47	36	29	7	9
80	( 788 )	( 575 )	( 11 )	( 19 )	( 27 )	( 53 )	( 43 )	( 32 )	( 7 )	( 10 )
	1073	637	12	12	44	53	51	37	8	10
	935	633	13	13	32	45	50	38	9	9
100	( 1004 )	( 635 )	( 12 )	( 13 )	( 38 )	( 49 )	( 51 )	( 38 )	( 9 )	( 10 )
	1222	807	9	10	53	69	64	46	9	10
	1257	860	12	20	40	81	67	44	12	9
125	( 1240 )	( 834 )	( 11 )	( 15 )	( 47 )	( 75 )	( 66 )	( 45 )	( 11 )	( 10 )
	1383*	1090	15	13	90	108	75*	59	15*	20
	1133*	1098	16	16	72	101	55*	84	15*	12
250	( 1258* )	( 1094* )	( 16 )	( 15 )	( 81 )	( 105 )	( 65* )	( 72 )	( 15* )	( 16 )
	488*	437*	9*	12*	44*	113	0*	92*	0*	18*
	378*	415*	14*	12*	34*	129	0*	58*	0*	14*
500	( 433* )	( 426* )	( 12* )	( 12* )	( 39* )	( 121 )	( 0* )	( 75* )	( 0* )	( 16* )
	0*	341*	0*	13*	0*	46*	0*	0*	0*	16*
	0*	339*	0*	13*	0*	37*	0*	0*	0*	7*
1000	0* )	( 340* )	( 0* )	( 13* )	( 0* )	( 42* )	( 0* )	( 0* )	( 0* )	( 12* )
Judgement	+	+	+	+	+	+	+	+	-	-
Specific Mutagenicity	8500	5180	36.0	167	212	368	304	176		
Positive Control	AF2	2AA	NaN <sub>3</sub>	2AA	AF2	2AA 20	AF2	2AA	9AA	2AA
Control	( 729 )	( 727 )	( 289 )	( 184 )	( 208 )	( 1262 )	( 363 )	( 394 )	( 660 )	( 131 )