

p-Chlorobenzyl chloride (p-クロロベンジルクロリド)

**Chemical Name:** p-Chlorobenzyl chloride  
**Synonym** 1-Chloro-4-(chloromethyl)benzene  
 Benzene, 1-chloro-4-(chloromethyl)-  
**Molecular weight:** 161.03  
**Melting point:** 28 - 30°C  
**Boiling point:** 213~214°C (760mmHg), 216-222°C  
**Flashing point:** 97 °C  
**Chemical Structure**

Clc1ccc(CCl)cc1

CAS No : 104-83-6  
 MITI No: (3)-78, (3)-91  
 M L No : 4-(15)-26, 4-(15)-37, 4-(15)-60  
 Source of Substance: Tokyo Kasei Kogyo Co. Ltd  
 Lot. No. : AX01  
 Purity : 98 %  
 Vehicle : DMSO

Experimental Data

Con. μg/ plate	Number of Revertants/plate									
	Base-substitution						Frame-shift			
	TA100		TA1535		WP2uvrA		TA98		TA1537	
	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
DMSO	(139)	(152)	(12)	(11)	(31)	(40)	(12)	(20)	(15)	(18)
	139	156	8	8	31	36	10	23	16	14
0.0763	(139)	(171)	(9)	(8)	(34)	(37)	(14)	(23)	(17)	(16)
	148	164	17	6	45	47	17	20	20	21
	135	165	8	13	46	32	20	24	14	23
0.305	(142)	(165)	(13)	(10)	(46)	(40)	(19)	(22)	(17)	(22)
	148	180	14	11	44	41	24	23	18	21
	179	151	9	8	38	47	22	26	17	22
1.22	(164)	(166)	(12)	(10)	(41)	(44)	(23)	(25)	(18)	(22)
	162	172	11	9	32	38	17	23	18	26
	153	166	11	6	33	33	23	32	18	23
4.88	(158)	(169)	(11)	(8)	(33)	(36)	(20)	(28)	(18)	(25)
	185	173	16	17	33	47	22	21	16	16
	178	195	9	11	45	39	20	30	22	22
19.5	(182)	(184)	(13)	(14)	(39)	(43)	(21)	(26)	(19)	(19)
	193*	241	10*	17	40	44	10*	26	13*	17
	205*	246	10*	10	57	56	13*	29	16*	18
78.1	(199*)	(244)	(10*)	(14)	(49)	(50)	(12*)	(28)	(15*)	(18)
	0*	0*	0*	0*	0*	0*	0*	0*	0*	0*
	0*	0*	0*	0*	0*	0*	0*	0*	0*	0*
IARC Evaluation : not yet cited	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)
	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)
	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)
	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)	(0*)
Judgement	-	-	-	-	-	-	-	-	-	-
Specific Mutagenicity										
Positive Control	AF2	2AA	NaN3	2AA	AF2	2AA	AF2	2AA	9AA	2AA
	(790)	(1212)	(240)	(247)	(242)	(724)	(396)	(359)	(412)	(183)

Experimental Data

Con. μg/ plate	Number of Revertants/plate									
	Base-substitution						Frame-shift			
	TA100		TA1535		WP2uvrA		TA98		TA1537	
	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
DMSO	(121)	(126)	(11)	(14)	(20)	(24)	(15)	(21)	(10)	(10)
	120		13				13		8	
	136		8				16		6	
1.22	(128)		(11)				(15)		(7)	
	128		7				16		10	
	120		7				22		7	
2.44	(124)		(7)				(19)		(9)	
	121	111	11	10	25	25	10	16	10	7
	127	109	16	10	17	30	10	13	13	10
4.88	(124)	(110)	(14)	(10)	(21)	(28)	(10)	(15)	(12)	(9)
	135	129	10	9	20	20	16	21	9	10
	138	115	7	11	25	25	14	25	6	11
9.77	(137)	(122)	(9)	(10)	(23)	(23)	(15)	(23)	(8)	(11)
	146	145	13	10	15	38	22	23	7	8
	155	142	13	13	26	23	17	30	6	7
19.5	(151)	(144)	(13)	(12)	(21)	(31)	(20)	(27)	(7)	(8)
	202*	149	16*	9	22	23	16*	16	10*	9
	146*	159	17*	13	28	28	23*	30	7*	10
39.1	(174*)	(154)	(17*)	(11)	(25)	(26)	(20*)	(27)	(9*)	(10)
	148*	207	6*	10	26	22	16*	31	7*	8
	155*	204	10*	8	30	38	16*	30	1*	7
78.1	(152*)	(206)	(8*)	(9)	(28)	(30)	(16*)	(31)	(4*)	(8)
		94*		13*	10*	37*		8*		9*
		112*		15*	2*	38*		15*		11*
156		(103*)		(14*)	(6*)	(38*)		(12*)		(10*)
		0*		0*	0*	0*		0*		0*
		0*		0*	0*	0*		0*		0*
313		(0*)		(0*)	(0*)	(0*)		(0*)		(0*)
Judgement	-	-	-	-	-	-	-	-	-	-
Specific Mutagenicity										
Positive	AF2	2AA	NaN3	2AA	AF2	2AA	AF2	2AA	9AA	2AA
Control	(942)	(1209)	(292)	(260)	(259)	(794)	(365)	(345)	(402)	(232)

Experimental Data

Con. $\mu$ g/ plate	Number of Revertants/plate					
	Base-substitution					
	TA102		TA104		WP2uvrA/pKM101	
	S9-	S9+	S9-	S9+	S9-	S9+
<u>DMSO</u>	( 297 )	( 363 )	( 339 )	( 358 )	( 204 )	( 268 )
	250	347	300	344	193	246
	292	344	290	324	197	256
<u>0. 0763</u>	( 271 )	( 346 )	( 295 )	( 334 )	( 195 )	( 251 )
	288	341	328	361	209	260
	250	361	301	389	195	262
<u>0. 305</u>	( 269 )	( 351 )	( 315 )	( 375 )	( 202 )	( 261 )
	305	362	340	344	245	261
	294	387	348	366	205	266
<u>1. 22</u>	( 300 )	( 375 )	( 344 )	( 355 )	( 225 )	( 264 )
	312	356	417	402	242	303
	310	380	404	439	243	271
<u>4. 88</u>	( 311 )	( 368 )	( 411 )	( 421 )	( 243 )	( 287 )
	294*	405	558*	493	347	433
	301*	358	527*	455	323	434
<u>19. 5</u>	( 298* )	( 382 )	( 543* )	( 474 )	( 335 )	( 434 )
	0*	201*	0*	372*	115*	295*
	0*	213*	0*	347*	129*	318*
<u>78. 1</u>	( 0* )	( 207* )	( 0* )	( 360* )	( 122* )	( 307* )
<u>313</u>	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )
<u>1250</u>	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )
<u>5000</u>	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )	( 0* )
Judgement	-	-	-	-	-	-
Specific Mutagenicity						
Positive	BLM	2AA	PA	2AA	AF-2	2AA
Control	( 959 )	(2584 )	(1780 )	(1072 )	(2991 )	(1067 )

Experimental Data

Con. μg/ plate	Number of Revertants/plate					
	Base-substitution					
	TA102		TA104		WP2uvrA/pKM101	
	S9-	S9+	S9-	S9+	S9-	S9+
DMSO	( 245 )	( 313 )	( 264 )	( 350 )	( 162 )	( 241 )
	246		278			
	216		265			
0.610	( 231 )		( 272 )			
	219		253			
	239		213			
1.22	( 229 )		( 233 )			
	227	307	274	332	139	256
	208	263	279	337	145	233
2.44	( 218 )	( 285 )	( 277 )	( 335 )	( 142 )	( 245 )
	248	319	280	334	163	207
	247	284	282	374	183	234
4.88	( 248 )	( 302 )	( 281 )	( 354 )	( 173 )	( 221 )
	240	320	291	368	178	246
	248	336	272	335	159	242
9.77	( 244 )	( 328 )	( 282 )	( 352 )	( 169 )	( 244 )
	236*	330	329	370	219	299
	261*	316	338	395	178	284
19.5	( 249* )	( 323 )	( 334 )	( 383 )	( 199 )	( 292 )
	243*	300	172*	397	239	317
	252*	313	390*	394	253	325
39.1	( 248* )	( 307 )	( 281* )	( 396 )	( 246 )	( 321 )
		293		447	248*	368
		325		416	304*	348
78.1		( 309 )		( 432 )	( 276 )	( 358 )
		252*		413*	96*	399*
		253*		462*	107*	423*
156		( 253* )		( 438* )	( 102* )	( 411* )
Judgement	-	-	-	-	-	-
Specific Mutagenicity						
Positive	BLM	2AA	PA	2AA	AF-2	2AA
Control	( 766 )	( 2053 )	( 1259 )	( 994 )	( 2205 )	( 1230 )

Experimental Data

Con. $\mu$ g/ plate	Number of Revertants/plate			
	Base-substitution			
	TA100		WP2uvrA/pKM101	
	S9-	S9+	S9-	S9+
<u>DMSO</u>	( 121 )	( 146 )	( 192 )	( 246 )
	150	130	216	252
	153	139	201	250
<u>12.5</u>	( 152 )	( 135 )	( 209 )	( 251 )
	185	121	246	319
	170	146	247	290
<u>25</u>	( 178 )	( 134 )	( 247 )	( 305 )
	190	198	318	391
	220	185	320	405
<u>50</u>	( 205 )	( 192 )	( 319 )	( 398 )
	171*	216	226*	463
	139*	207	255*	494
<u>75</u>	( 155* )	( 212 )	( 241* )	( 479 )
	0*	197	117*	477
	0*	253	126*	521
<u>100</u>	( 0* )	( 225 )	( 122* )	( 499 )
	0*	193*	0*	314*
	0*	171*	0*	301*
<u>125</u>	( 0* )	( 182* )	( 0* )	( 308* )
	0*	300	0*	164*
	0*	313	0*	204*
<u>150</u>	( 0* )	( 307 )	( 0* )	( 184* )
		293	0*	145*
		325	0*	131*
<u>175</u>		( 309 )	( 0* )	( 138* )
		252*	0*	0*
		253*	0*	0*
<u>200</u>		( 253* )	( 0* )	( 0* )
Judgement	-	-	-	-
Specific Mutagenicity				
<u>Positive Control</u>	AF2	2AA	AF2	2AA
	( 945 )	(1238 )	(2666 )	(1166 )