

Mutations in BxPC-3

BXPC3_PANCREAS													
Gene	Protein Change	Chr	Start	End	NCBI Build	Variant Classification	Variant Type	Reference Allele	Tumor Allele 1	dbSNP ID	Annotation Transcript	Other Transcripts	Mutation Assessor
AAK1	p.S541_542QQ>C	2	69741754	69741756	37	In_Frame_Del	Del	TGT	-	rs55712143	uc002sfp.2	AK1_uc010fdk.2...	
ADAM17		2	9630245	9630245	37	3'UTR	Del	T	-		uc002qzu.2	AH1_uc010yiz.1...	
ADAM28		8	24211994	24211997	37	3'UTR	Del	CTAT	-	rs10610134	uc003xdy.2	AM28_uc011laa...	
AKAP12	p.L1531_1532ins	6	151674116	151674117	37	In_Frame_Ins	Ins	-	GAG	rs34737819	uc011eep.1	AP12_uc003qoe...	
AKAP6	p.T80S	14	32902937	32902937	37	missense_Mutati	SNP	A	T		uc001wrq.2	AKAP6_uc010aml	view
ALPK2	p.E210T	18	56204388	56204402	37	In_Frame_Del	Del	CAGTT...	-	rs44017;rs11762	uc002lhj.3	LPK2_uc002lhk...	
BCL11B		14	99640466	99640466	37	3'UTR	SNP	C	G		uc001yga.2	L11B_uc001ygl	view
BMP4		14	54416601	54416602	37	3'UTR	Ins	-	AT	495140;rs76331	uc001xal.3	BP4_uc010aoh.2...	
CACNB2		10	18828669	18828670	37	3'UTR	Ins	-	C	022725;rs58831	uc001ipr.2	CACNB2_uc009xjz...	
CDK10	p.S293L	16	89761424	89761424	37	missense_Mutati	SNP	C	T		uc010cio.2	CDK10_uc002fod	view
CDK18		1	205495890	205495891	37	Intron	Del	CT	-		uc001hcr.2	CDK18_uc010pri...	
CHD1	p.P1684del	5	98192165	98192167	37	In_Frame_Del	Del	AGG	-	267787;rs61749	uc003knf.2	CHD1_uc010jbn.2...	
CLTCL1	p.V1201_splice	22	19189003	19189004	37	Splice_Site_Ins	Ins	-	C	rs11386977	uc002zpb.2	CLTCL1_uc011agv...	
CNTN6		3	1339681	1339686	37	Intron	Del	GTTTT...	-		uc003boz.2	CNTN6_uc010hbo...	
CREB3L2	p.T100del	7	137612914	137612916	37	In_Frame_Del	Del	TGG	-		uc003vtw.2	CREB3L2_uc003vt...	
DBF4	p.D82G	7	87514319	87514319	37	missense_Mutati	SNP	A	G		uc003ujf.1	DBF4_uc003ujh.1	view
DMPK	p.V173I	19	46281893	46281893	37	missense_Mutati	SNP	C	T		uc002pdi.1	DMPK_uc010xss.1	view
DYRK1A		21	38884966	38884967	37	3'UTR	Ins	-	T	rs72430212	uc002ywk.2	DYRK1A_uc002ywi...	
EDN1		6	12290729	12290729	37	5'UTR	Del	A	-		uc003nae.3	EDN1_uc010jpb.2...	
EP300	p.R397*	22	41525914	41525914	37	missense_Mutati	SNP	C	T		uc003azl.3		view
EPHA6		3	97367230	97367231	37	Intron	Ins	-	A	rs115253553	uc010how.1	EPHA6_uc003drs...	
ERCC6	p.L422F	10	50732212	50732212	37	missense_Mutati	SNP	T	A		uc001jhs.3	ERCC6_uc001jht	view
ETV1		7	14028708	14028708	37	5'UTR	Del	A	-	rs77190713	uc011jq.1	ETV1_uc011jxn.1...	
EXT1		8	118811807	118811807	37	3'UTR	Del	T	-		uc003yok.1		
FBN1	p.A979T	15	48782195	48782195	37	missense_Mutati	SNP	C	T		uc001zwx.1		view
FGFR3		4	1809128	1809129	37	3'UTR	Del	GT	-		uc003gdu.2	FGFR3_uc003gdr...	
FLI1		11	128638217	128638217	37	Intron	Del	G	-		uc010sbu.1	FLI1_uc010sbt.1...	
GPR112	p.D2657del	X	135474445	135474447	37	In_Frame_Del	Del	GAT	-	rs3030295	uc004ezu.1	GPR112_uc010nsb...	
GRIA3		X	122318386	122318387	37	Splice_Site_Ins	Ins	-	G	rs66632982	uc004etq.3	GRIA3_uc004etr...	
GUCY2C		12	14827464	14827466	37	Intron	Del	ATC	-	rs525754;rs3083	uc001rcd.2	GUCY2C_uc009zhz...	
HNF1A		12	121434630	121434631	37	Intron	Ins	-	TCATTCAT	rs75100977	uc010sxn.1	HNF1A_uc001tze...	
HNRNP11		5	179050364	179050364	37	Intron	SNP	G	C		uc003mkf.3	HNRNP11_uc003n	view
ILK		11	6630029	6630029	37	Intron	Del	C	-		uc001mee.2	ILK_uc001mef.2...	

INHBA	p.Y404S	7	41729318	41729318	37	issense_Mutati	SNP	T	G	rs113772875	uc003thq.2	IHBA_uc003thr.	view
INSRR	p.G331V	1	156821530	156821530	37	issense_Mutati	SNP	C	A		uc010pht.1	TRK1_uc001fqf.	view
ITPR2	p.R470_splice	12	26834804	26834805	37	Splice_Site_Ins	Ins	-	ACTC	696203;rs59199	uc001rhg.2		
JARID2	p.K240E	6	15487585	15487585	37	issense_Mutati	SNP	A	G		uc003nbj.2	JRID2_uc011diu	view
KLK7		19	51480672	51480675	37	3'UTR	Del	GGTT	-		uc002puo.2	KLK7_uc002pup.2...	
MAF		16	79628325	79628325	37	3'UTR	Del	A	-		uc002ffm.2	MAF_uc002ffn.2_...	
MAML3	p.Q767_splice	4	140651585	140651587	37	Splice_Site_Del	Del	CTG	-		uc003ihz.1	GST2_uc010ioi....	
MAP3K1	p.T949del	5	56177849	56177851	37	In_Frame_Del	Del	CAA	-		uc003jqw.3		
MAP3K14	p.R218_splice	17	43364293	43364294	37	Splice_Site_Ins	Ins	-	G	rs66766023	uc002iiv.1	MAP3K14_uc010da...	
MYT1	p.R393L	20	62839727	62839727	37	issense_Mutati	SNP	G	T		uc002yii.2	MYT1_uc002yih.2	view
NCAM1		11	112832307	112832308	37	5'UTR	Ins	-	A	rs117108942	uc009yyq.1	NCAM1_uc001pno....	
NCAM1		11	112832340	112832340	37	5'UTR	Del	C	-		uc009yyq.1	NCAM1_uc001pno....	
NCAM1	p.K161E	11	113078643	113078643	37	issense_Mutati	SNP	A	G		uc009yyq.1	NCAM1_uc001pnc	view
NCOA3	p.QQQ1272del	20	46279815	46279823	37	In_Frame_Del	Del	GCAGC...	-		uc002xtk.2	NCOA3_uc010ght....	
NPR1		1	153665919	153665920	37	3'UTR	Ins	-	C	rs66682713	uc001fcs.3	NPR1_uc010pdz.1...	
NR1H2	p.176_177insQ	19	50881820	50881821	37	In_Frame_Ins	Ins	-	AAC	rs783852;rs11235	uc010enw.2	NR1H2_uc002prv....	
NR3C1		5	142780419	142780419	37	Splice_Site_SNP	SNP	T	A		uc003lmy.2	NR3C1_uc003lmz	view
NUP214		9	134067556	134067557	37	Intron	Ins	-	T	rs66652901	uc004cag.2	NUP214_uc004cah...	
PAX3		2	223158653	223158653	37	Intron	Del	T	-	rs112753212	uc002vmt.1	PAX3_uc002vmy.1...	
PDE4DIP	p.Q1665fs	1	144873963	144873963	37	In_Frame_Shift_Del	Del	T	-		uc001elw.3	PDE4DIP_uc009wir...	
PDE4DIP	p.V486fs	1	144917828	144917828	37	In_Frame_Shift_Del	Del	A	-		uc001elw.3	PDE4DIP_uc009wir...	
PDE4DIP		1	144931366	144931367	37	Intron	DNP	CC	TA		uc001elw.3	PDE4DIP_uc009wir...	
PIK3C2G	p.P129del	12	18435399	18435401	37	In_Frame_Del	Del	CCC	-		uc010sib.1	PIK3C2G_uc001rd...	
PIK3C2G		12	18800990	18800990	37	3'UTR	Del	A	-	rs16914368	uc010sib.1	PIK3C2G_uc001rd...	
PLK2		5	57754052	57754052	37	Intron	SNP	A	G		uc003jrn.2	PLK2_uc011cql.1	view
PMS1		2	190670539	190670540	37	Intron	Ins	-	A	rs3214425	uc002urh.3	PMS1_uc010zga.1...	
PPP3R1	p.F15_splice	2	68415824	68415824	37	Splice_Site_SNP	SNP	T	A		uc002sei.1		view
PRKDC	p.L1244fs	8	48805816	48805817	37	In_Frame_Shift_Ins	Ins	-	G	rs67588121	uc003xqi.2	PRKDC_uc003xqi....	
RECQL4	p.R766fs	8	145738768	145738768	37	In_Frame_Shift_Del	Del	G	-		uc003zjd.2		
REM1		20	30064232	30064233	37	5'UTR	Ins	-	C		uc002wwa.2		
RNASEL	p.M641T	1	182545508	182545508	37	issense_Mutati	SNP	A	G		uc001gpj.1	RNASEL_uc009wx	view
SLC45A3	p.D547N	1	205628385	205628385	37	issense_Mutati	SNP	C	T		uc001hda.1	SLC45A3_uc010p	view
TEC		4	48139307	48139310	37	3'UTR	Del	TTAT	-	rs79098351	uc003gxz.2		
TP53	p.Y220C	17	7578190	7578190	37	issense_Mutati	SNP	T	C		uc002gim.2	TP53_uc002gig.1	view
TRPS1		8	116635872	116635873	37	5'UTR	Ins	-	A	rs76893349	uc003yny.2	TRPS1_uc011lhy....	
TTN		2	179391639	179391640	37	3'UTR	Ins	-	T	rs36037175	uc010zfg.1	TTN_uc010zfh.1_...	
TTN	p.R14123T	2	179477180	179477180	37	issense_Mutati	SNP	C	G		uc010zfg.1	TTN_uc010zfh.1_...	view
TTN	p.T799M	2	179650444	179650444	37	issense_Mutati	SNP	G	A		uc010zfg.1	TTN_uc010zfh.1_...	view

VEGFC	p.S419_splice	4	177605082	177605084	37	Splice_Site_Del	Del	TCA	-		uc003ius.1	
VPS13B		8	100887922	100887923	37	3'UTR	Del	TG	-		uc003yiv.2	'S13B_uc003yiw...
YSK4	p.L212L	2	135745806	135745806	37	Silent	SNP	C	T		uc002tue.1	SK4_uc002tuf.1 view

