

## Key

### mutation

- deletion loss of genetic material
- DR down regulation; decrease mRNA transcripts
- insertion gain of genetic material (eg duplication, amplification, transposition)
- UR up regulation; increase mRNA transcripts
- AE altered expression (not stated whether up or down)
- LOH loss of heterogeneity
- OE over expression related to experimental factors

### Grade

- TNM
- GS; gleason score
- GG; gleason grade
- BPH; benign prostatic hyperplasia
- NORM; normal prostate tissue
- HR; hormonal refractory prostate
- PIN; prostate neoplasm
- Prim; primary
- Recur; recurrent
- Mets; metastases
- Local; localised prostate tumor
- Cell Lines: PC3, DU145, LNCaP
- TURP;trans urethral prostate chips

### Technique

- CGH; comparative genomic hybridisation
- FISH; fluorescent *in situ* hybridisation
- GC; gene chips, expression microarrays
- TMA; tissue microarrays
- IH; immunohistochemistry
- WB; western blot
- SB; southern blot
- NB; norther blot
- PCR; polymerase chain reaction
- RT-PCR; reverse transcriptase PCR
- MSI; microsatellite instability
- MSA; microsatellite amplification

Loci	mutation	possible gene of interest	possible role in Prostate cancer	Grade	Technique	%	ref
<b>Chromosome 1</b>							
1p	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH & FISH	54	7
1p13		KREV-1, ras related gene	Putative Tumor Suppressor Gene				4
1p13.2	DR	GSTM3( glutathione transferase M3)		GG4/5, BPH,	GC	100	8
1p13.3	AE	RAP1A ( RAS oncogene family).		GS5-10, BPH	GC		11
	AE	GSTM1 (glutathione S-transferase M1)		GS5-10, BPH,	GC		11
1p31	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH & FISH	Minimal	7
1p34.3	UR	MACMARCKS: macrophage myristoylated alanine-rich C kinase substrate.		GG4/5, BPH,	GC	100	8
	UR	MACMARCKS		GS5-9, norm	GC		12
1p34.1	UR	TSPAN1; tetraspan 1					
1p35	AE	Splicing factor, arginine/serine rich 4 (?)		GS5-10, BPH	GC		11
1p35.2-p36.1	AE	Gardner-Rasheed feline sarcoma viral (v-fgr) homolog	oncogene	GS5-10, BPH	GC		11
1p36-pter	deletion	Contains p53-related putative neuroblastoma TSG		T1-4,Nx,Mx,0,1 vs norm	CGH & FISH	Minimal	7
1p36.1-35	AE	Solute carrier family 9 (sodium hydrogen exchanger), isoform 1		GS5-10, BPH	GC		11
1p36.1	AE	Elastase 3, pancreatic (protease E)		GS5-10, BPH	GC		11
	AE	ECE1 endothelin converting enzyme 1		GS5-10, BPH	GC		11
1p36.2	DR	SLC2A5 (solute carrier family 2, member 5)		GG4/5, BPH,	GC	100	8
	AE	VAM3 (vesicle-associated membrane protein 3 (cellubrevin)		GS5-10, BPH	GC		11
1p36.3	LOH	PKCz very distal;			FISH		4
	AE	CDC10 (cell division cycle 10, homolog).		GS5-10, BPH,	GC		11
	AE	Rho guanine exchange factor 16	Putative neuroblastoma protein,	GS5-10, BPH	GC		11
	AE	KIAA0720		GS5-10, BPH,	GC		11
1q	insertion			GS5-10	CGH	16	1
1q21	AE	Thrombospondin		GS5-10, BPH,	GC	(2/15)	11
	AE	3Aryl hydrocarbon receptor nuclear translocator		GS5-10, BPH,	GC		11
	insertion		possible oncogene	GS5-10	CGH		1
	insertion			GS5-10	CGH		1
1q21.3	deletion	nm-23 H1 antioncogene		GS 8-10, stage D	IH		23
1q22	AE	Phosphoprotein enriched in astrocytes 15		GS5-10, BPH	GC		11
	AE	Ric (Drosophilla)-like expressed in many tissues		GS5-10, BPH	GC		11
	AE	Coatomer protein complex, a, (?)		GS5-10, BPH	GC		11
1q24	AE	Quiescin Q6		GS5-10, BPH	GC		11
1q25		RNASEL: 2'-5'-oligoadenylate (2-5A) dependent RN ase L	Proposed Tumor Suppressor Gene			9, 53	
	insertion			GS5-10	CGH	(1/15)	1
1q25.1-q32.3	AE	Ladinin1; LAD1, 120kDa linear IgA bullous dermatosis antigen		GS5-10, BPH	GC		11
1q25-q32	insertion			T1-4,Nx,Mx,0,1 v norm	CGH & FISH		7
1q31-q32	AE	Protein tyrosine phosphatase, receptor type C		GS5-10, BPH	GC		11
1q43	AE	Nidogen (enactin; NID)		GS5-10, BPH	GC		11

**chromosome 2**

2p13	DR	ANNXA 4, annexin 4		GS6-9, BPH, HR	GC		6
2p15.2	AE	Catenin (cadherin-associated protein), delta 2		GS5-10, BPH	GC		11
2p21	LOH	PKCe			FISH	100	4
	AE	3-hydroxyanthranilate 3,4-dioxygenase		GS5-10, BPH,	GC		11
	AE	endothelial PAS domain protein 1 (homolog).		GS5-10,BPH	GC		11
	UR	TACSTD1: tumor associated calcium signal transducer 1		GS5-9, norm	GC		12
2p22-p21	AE	Latent transforming growth factor beta binding protein 1		GS5-10, BPH	GC		11
2p22.3-p21	DR	BRF2(butyrate response factor 2		GG4/5, BPH	GC	100	8
2p23	AE	KCNK3; potassium channel, subfamily K, member 3 (TASK)		GS5-10, BPH	GC		11
2p23-24		NMYC; .	possible connection with MYC	Prim, Recur, Mets vs BPH	FISH, TMA		13
2p25	AE	Tumor suppressing subtransferable candidate 1 (TSSC1)		GS5-10, BPH	GC		11
	UR	AF052107		GS5-9, norm	GC		12
2q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH & FISH	27	7
	deletion			pT2N0	CGH	18	22
	deletion			GS5-10	CGH		1
2q12-q14	DR	DRAL (down-regulated in rhabdomyosarcoma lim protein)		GG4/5, BPH	GC	100	8
2q21-q22	deletion			GS5-10	CGH		1
2q23.3	AE	Ras homolog gene family, member E		GS5-10, BPH	GC		11
2q31-32.1	AE	CHN1 (chimaerin)		GS5-10, BPH	GC		11
2q33	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH & FISH		7
	UR	PLCE (phospholipase C, epsilon)		GG4/5, BPH,	GC	100	8
	deletion	AOX1 (aldehyde oxidase 1)		GG4/5, BPH,	CGH	100	1
2q35	DR	DES (desmin)		GG4/5, BPH,	GC	100	8
2q37	AE	Collagen, type VI, alpha 2		GS5-10, BPH,	GC		11
<b>Chromosome 3</b>							
C3	deletion		deletion only in hormone Refractory	T1-4,Nx,Mx,0,1 vs norm	CGH & FISH	100	7
3p25-26	LOH	includes the <i>von Hippel Lindau gene</i>		GS5-9, T2& T3N0M0, TXN+M0	MSI	20	10
3p24-p22	DR	RAB5A, member of the RAS oncogene family		PCa, BPH	GC		5
3p21.2-p14.3	AE	Sarcolemmal-associated protein		GS5-10, BPH	GC		11
3p21.1	UR	Arginine-rich protein		GG 4/5, BPH,	GC	100	8
	AE	Arginine-rich protein		GS5-10, BPH,	GC	1	11
3p21.2	LOH	PKCd?			FISH		4
3p21.3	AE	Chemokine (C-X3-C)receptor 1		GS5-10, BPH,	GC		11
3cen-q21	DR	MYLK (myosin, light polypeptide kinase)		GG4/5, BPH,	GC	100	8
	DR	CSTA (cystatin A or stefin A) @ 3q21		GG4/5, BPH,	GC	100	8
3q	insertion			T1-4,Nx,Mx,0,1 vs BPH	CGH& FISH	24	7
	insertion			GS5-10	CGH	18	1
3q13.1-13.2	UR	ALCAM/activated leucocyte cell adhesion molecule		GG4/5, BPH,	GC	100	8
3q21	AE	CSRP2 (cysteine and glycine-rich protein)	possible oncogene	GS5-10, BPH,	GC	(1/15)	11
	insertion			GS5-10	CGH		1
	insertion			GS5-10	CGH		1
3q21-q23	AE	EphB1		GS5-10, BPH	GC		11
3q22-q24	DR	TRPC1 (transient receptor potential channel 1)		GG4/5, BPH,	GC	100	8
3q25.1-26.1	AE	SWI/SNF related		GS5-10, BPH	GC		11
3q26	LOH	PKCi;?			FISH		4
3q25-26	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
3q26.1	DR	HKvBeta3 (potassium voltage-gated channel, b member 3)		GG4/5, BPH,	GC	100	8
3q 28	DR	ETV5 (ets variant gene 5[ets-related molecule])		GG4/5, BPH,	GC	100	8
	AE	ETV5		GS5-10, BPH,	GC		11
	OE	ETV5	possible response to ischemia	Radical prostatectomy tissue	GC, WB, NB, IH		14
3q29	AE	3-hydroxybutyrate dehydrogenase (heart, mitochondria)		GS5-10, BPH	GC		11

**Chromosome 4**

4q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	19	7
	deletion			GS5-10	CGH	18	1
4q12	AE	Sarcoglycan, beta		GS5-10, BPH	GC		11
4q13-q23	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
4q21	AE	Small inducible cytokine B subfamily		GS5-10	GC		11
	insertion		Found in metastases only	GS5-10	CGH	(1/15)	1
4q22	AE	LIM protein		GS5-10, BPH	GC		11
	AE	LIM		Prostate cancer, BPH	GC		5
	UR	AL049969		GS5-9,norm	GC		12
4q22.1	AE	HEVIN; SPARC-like 1 (mast 9, hev1)		Prostate cancer, BPH	GC		5
4q26-q27	deletion		possible tumor suppressor gene	GS5-10	CGH		1
4q28.3	insertion		Found in metastases only	GS5-10	CGH	(1/15)	1
4q34-q35	DR	FAT tumor suppressor homolog 1 (drosophila)		Metastases	GC		5
<b>chromosome 5</b>							
5p	deletion			T2N0	CGH		22
5p13.2-11.1	AE	AMACR; alpha-methylacyl-CoA racemase		GS5-10, BPH	GC		11
	UR	AMACR		GS5-9, norm	GC		12
5q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	14	7
	deletion			T1N0M0	CGH	4.2	21
	deletion			T2N0	CGH		22
	deletion			GS5-10	CGH	32	1
5q2	UR	HSD17B4/17b- hydroxysteroid dehydrogenase IV		GG4/5, BPH,	GC	100	8
5q11	deletion		possible Tumor Suppressor Gene.	T2N0	CGH	6	22
5q12-23	LOH	eg APC		GS5-9, T2& T3N0M0, TXN+M0	MSI	17	10
5q13.1	UR	Neuronal apoptosis inhibitory protein		GG4/5, BPH,	GC	100	8
5q13	AE	MAP1B; microtubule-associated protein 1B		GS5-10, BPH	GC		11
5q15-q21	deletion	Contains the APC/MCC region	possible Tumor Suppressor Gene.	GS5-10	CGH		1
5q15-q23	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
5q14-q31	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
5q23	AE	Sorting nexin 2 (SNX2)		GS5-10, BPH	GC		11
5q31.1	AE	ETTF1; euk translation termination factor 1		GS5-10, BPH	GC		11
5q32-q34	DR+B171	ANXA 6, annexin VI (p68)		GG4/5, BPH,	GC	100	8
	AE	annexin VI (p68)		GS5-10, BPH,	GC		11
	DR	DPYSL3 (dihydropyrimidinase-like 3)		GG4/5, BPH,	GC	100	8
5q32-33	AE	Protocadherin 1 (cadherin-like 1)		GS5-10, BPH	GC		11
5q34	DR	KCNMB1( potassium large conductance calcium activated channel)		GG4/5, BPH,	GC	100	8
	AE	KCNMB1		GS5-10, BPH,	GC		11
5q35.3	AE	Enigma (LIM domain protein)		GS5-10, BPH	GC		11

**chromosome 6**

6p21.2	deletion	(p21 WAF1/CIP1 cyclin kinase inhibitor: can induce G1 arrest						
6p21	insertion		possible Tumor Suppressor Gene.	GS5-10	CGH			1
6p21.2	UR	PIM1, pim1 oncogene	regulation tied to MYC	Mets, local, BPH, PIN, norm	GC			5
6p21.31	AE	GABA B receptor 1(gamma-aminobutyric acid		GS5-10, BPH	GC			11
6p22-p21	DR	ID4 (inhibitor of DNA binding 4)		GG4/5, BPH,	GC	100		8
6q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	27		7
	deletion			T2N0	CGH			22
	deletion			GS5-10	CGH	43		1
	LOH			GS5-9, T2& T3N0M0, TXN+M0	MSI	24		10
6q15-q16	deletion		possible Tumor Suppressor Gene.	GS5-10	CGH			1
6q16	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH			7
6q12-q21	AE	Glycoprotein hormones, alpha polypeptide chain precursor. Or TAK1; transforming growth factor beta-activated kinase 1		GS5-10, BPH,	GC			11
6q21	deletion	CCNC; encodes cell-cycle regulatory protein cyclin		T2N0	CGH			22
6p22	AE	Butyrophilin, subfamily 2, member A2 (BTN2A2)		GS5-10, BPH	GC			11
6q21-q23.2	DR	GJA1 (gap junction protein, a 1, 43kD(connexin 43))		GG4/5, BPH,	GC	100		8
6q24-qter	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH			7
6q25.1	AE	Estrogen receptor 1 (ESR1)		GS5-10, BPH	GC			11
<b>Chromosome 7</b>								
C7	insertion	Contains: CLK2, MDR1 and elongation factor-1-g.	Trisomy associated with progression	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH			7
	insertion							26
	DR	RARRES2(retinoic acid receptor responder 2)		GG4/5, BPH,	GC	100		8
7p	insertion			HR	CGH& FISH	32,0	7, 54	
7p11	AE	Dopa decarboxylase (aromatic L-amino acid decarboxylase)		GS5-10, BPH,	GC	100		11
7p15-14	UR	T-cell receptor Ti rearranged g-chain		GG4/5, BPH,	GC	100		8
7p15-p21	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH			7
7p22	AE	Islet cell autoantigen (69kDa)		GS5-10, BPH	GC			11
7q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	43		7
	insertion			T1NxMx	CGH	4.2		21
	insertion			GS5-10	CGH	21		1
7q11.2	insertion		possible Tumor Suppressor Gene.	GS5-10	CGH			1
7q11.21	AE	PDGF associated protein		GS5-10, BPH	GC			11
7q11.23	AE	Claudin 4, CPE receptor (clostridium perfringens enterotoxin)		GS5-10, BPH	GC			11
7q21	insertion	Contains HGF (hepatocyte growth factor)		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7, 55	
	OE	HGF	overexpressed in response to ischemia	radical prostatectomy tissue	GC, WB, NB, IH			14
	AE	Cytochrome P450 3A5 human;		GS5-10, BPH	GC			11
7q21-22.1	DR	CYP3A7- cytochrome P450 (P-450 HFLa)		GG4/5, BPH,	GC	100		8
7q22	AE	Procollagen C-endopeptidase enhancer		GS5-10, BPH,	GC			11
7q22-q31.1	AE	PRKAR2B; Protein kinase, cAMP-dependent, regulatory, type II		GS5-10, BPH	GC			11
7q31	insertion	Caveolin (D7S486), MET (receptor for HGF)		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	30100		7
	LOH	2 possible genes, (not MET);.	possible Tumor Suppressor Gene.	T3N0M0	PCR MSA			16
	DR	CAV (caveolin 1, caveolae protein, 22Kda)		GG4/5, BPH	GC			8
	insertion		possible Tumor Suppressor Gene.	GS5-10	CGH			1
7q31.1	DR	Caveolin 2		GG4/5, BPH,	GC	100		8
	AE	Caveolin 2		GS5-10, BPH,	GC			11
7q32	insertion		Found in metastases only	GS5-10	CGH		(1/15)	1
7q33	DR	HBNF-1 (nerve growth factor)		GG4/5, BPH,	GC	100		8
	AE	HSPCO49 protein		GS5-10, BPH,	GC			11
7q36-q35	AE	Potassium voltage-gated channel, subfamily H, member 2		GS5-10, BPH,	GC	100		11

**Chromosome 8**

8p	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	73	7
	deletion			T1NxMx	CGH	10.4	21
	deletion			T2NO	CGH		22
	deletion			GS5-10	CGH	>48	1
	LOH			GS5-9, T2& T3N0M0, TXN+M0	MSI	69	10
8p11.2-p11.1	deletion			N/A, 17 malignant tumors, 1 benign	CGH,FISH,PCR		2
8p12-p22	DR	FGFR1 (fibroblast growth factor receptor 1)		GG4/5, BPH,	GC	100	8
	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
		LPL					56
8p21	UR	TRPM-2(testosterone-repressed prostate message 2		GG4/5, BPH,	GC	100	8
	deletion	NKX3.1 human prostate specific, androgen-reg homeobox gene;		T2NO	CGH		22
	AE	NEFL (16)Tumor protein D52		GS5-10, BPH	GC		11
	deletion	NKX3.1	possible Tumor Suppressor Gene.	GS5-10	CGH		1
8p21.2	deletion			T2NO	CGH		22
8p22	deletion	Contains NKX3.1 gene: prostate specific,	possible Tumor Suppressor Gene.	T2NO	CGH	Common	22
	deletion	N33	possible Tumor Suppressor Gene.			rare	17
	deletion	LZTS1 (FEZ1)	possible Tumor Suppressor Gene.	T1NxMx	CGH		21
	deletion			N/A			57
	AE	human acid ceramidase gene;	promotes apoptosis	GS5-7, BPH, cell lines, norm	SB, NB, RT PCR	15/36	18
	deletion	LPL; lipoprotein lipase		GS5-9 (T3), Mets (N1)	SB,MSI	47	19
	deletion	MSR;macrophage scavenger receptor gene;		GS5-9 (T3), Mets (N1)	SB,MSI	69	19
8p 22-p21	All Imb		2 possible Tumor suppressor genes	T3NoMo	PCR MSA	58	16
8p22-pter	deletion		possible Tumor Suppressor Gene.	GS5-10	CGH		1
8p23	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
8p23.1-21.3	DR	IGFBP5 = NAT1 (N-acetyl transferase)		Metastases	GC		5
8q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	73	7
	insertion			T2NO	CGH		22
	deletion			T2NO	CGH		22
	insertion			GS5-10	CGH	36	1
	insertion	whole arm of q.		N/A, 17 malignant tumors, 1 benign	CGH,FISH,PCR		2
8q21	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
8q21.13	AE	E2F transcription factor 5, p130-binding		GS5-10, BPH,	GC	100	11
8q22.3-q23	DR	KIAA0003/ANGPT1 (angiopoietin 1)		GG4/5, BPH,	GC	100	8
8q23-qter	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH & FISH		7
8q24	insertion	Contains c-MYC: regn of cell proliferation, diff. and apoptosis.		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	10	7
	insertion	c-MYC	oncogene,	Prim, Recur, Mets, vs BPH,	FISH TMA	<11	13
	insertion	c-MYC	Protooncogene	HR			55
	insertion	c-MYC; gene amplications (low copy no.)	oncogene	GS5-10	CGH		1
	insertion	MYC proto-oncogene		GS4-10, BPH	FISH	33	15
8q24.1	UR	Ectonucleotide pyrophosphatase/phosphodiesterase 2		Prostate cancer, BPH	GC		5
8q24.12-q24.13	AE	v-myc avian myelocytomatosis viral oncogene homolog		GS5-10, BPH	GC		11

Chromosome 9							
9p21	deletion	p16 (INK4A/MTS1/CDKN2);	multiple tumor suppressor gene site	N/A, 32 Prostate Tumors, 2 cell lines	SB, PCR	(1 / 32)	20
9q	insertion			T1NxMx	CGH	6.3	21
	gains			T2NO	CGH		22
	LOH			GS5-7, 8&9, HR, Mets, norm	SB	15	25
	insertion			GS5-10	CGH	18	1
9q12-q21.2	DR	ANXA1, annexin 1		GS6-9, BPH, HR	GC		6
9q21-q22	insertion	PCA3 (DD3),non coding RNA	possible oncogene	T2NO	CGH		22
	AE	DD3;prostate specific		BPH, Mets, norm	NB, RT PCR	95	27
	DR	RNAROR2 (receptor tyrosine kinase-like orphan receptor 2)		GG4/5, BPH,	GC	100	8
9q21.3-22.1	DR	gas 1 gene		GG4/5, BPH,	GC	100	8
9q22	AE	CDC28 protein kinase 2 (CKS2)		GS5-10, BPH	GC		11
9q32-qter	insertion		possible oncogene	GS5-10	CGH		1
9q34.1	insertion	ABL	oncogene	T2NO	CGH		22
9q34.2-qter	insertion		Possible oncogene	T1NxMx	CGH		21
Chromosome 10							
10p	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	22	7
10p11	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
10p15		PKCq very distal,			FISH		4
10p22	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	22	7
10q	LOH		possible Tumor Suppressor Gene.	Local GS5-7, 8&9, HR, Mets,norm	SB	29	25
	deletion			GS5-10	CGH	18	1
10q11.1	DR	SDF1 (stromal cell-derived factor 1)		GG4/5, BPH,	GC	100	8
	AE	SDF1 (stromal cell-derived factor 1)		GS5-10, BPH,	GC		11
10cen-q21	deletion		tumor suppressor gene	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
10q21.1	AE	EGR2;early growth response 2 (Krox-20(drosophila homolog)		GS5-10, BPH	GC		11
10q21.1-21.2	DR	ANXA 7; annexin 7		GS6-9, BPH, HR	GC		6
10q21.3	insertion		Found in metastses only	GS5-10	CGH	(1/15)	1
10q22	DR	CAMK2G (calcium/calmodulin-dependant protein kinase		GG4/5, BPH,	GC	100	8
	AE	CAMK2G		GS5-10, BPH	GC		11
10q22.1-qter	deletion	Mxi1; candidate tumor suppressor gene		GS4-9, Lymph node mets, & mets, HR (GS6-10) vs TURP& blood	CGH, SB & PCR MSA		3
10q22.1-q23	DR	VCL (vinculin)		GG4/5, BPH,	GC	100	8
10q23	deletion	PTEN; protein tyrosine phosphatase gene	tumor suppressor gene	T2NO	CGH		22
	deletion	PTEN	tumor suppressor gene	cell lines;( LNCaP)	RT PCR, SB	100	28
	DR	PTEN		Prostate cancer	GC		5
	deletion	TEP1/PTEN/MMAC1;+C33		cell lines	NB		30
	DR	ANXA 11, annexin 11		GS6-9, BPH, HR	GC		6
10q23.3	deletion	MMAC1	possible Tumor Suppressor Gene.			26, 3	
10q23-q24	DR	RBP (retinal binding protein)		GG4/5, BPH,	GC	100	8
	AE	Retinal binding protein 4, interstitial		GS5-10, BPH,	GC		11
10q24	deletion	MXI1; a negative regulator of the MYC oncoprotein	possible Tumor Suppressor Gene.	GS5-10	CGH		1
	LOH	RB; retinoblastoma gene; suppress tumorigenicity		Prim, Mets, BPH	IH, PCR	10 to 30	32
10q25	deletion	MXI-1	possible Tumor Suppressor Gene.			Rare 26, 31	
	deletion	MXI-1	tumor suppressor gene	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	46	7
10q26	deletion		tumor suppressor gene	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
	DR	Fibroblast growth factor receptor K-Sam		GG4/5, BPH,	GC	100	8

<b>chromosome 11</b>								
11p11.2	UR	PSMA		GG4/5, BPH,	GC	100	8	
11p15.1	AE	RIG, regulated in glioma		Mets, local, BPH, PIN, norm	GC		5	
11p15.5	DR	H19 RNA gene		GG4/5, BPH,	GC	100	8	
	DR	IGF II (insulin-like growth factor 2)		GG4/5, BPH,	GC	100	8	
11p15.5-p15.3	AE	Lactate dehydrogenase C (LDHC)		GS5-10, BPH	GC		11	
11q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	22	7	
11q11	AE	PRO1659 protein		GS5-10, BPH	GC		11	
11q12	AE	Protein phosphatase 2, regulatory subunit B (B56) b isoform		GS5-10, BPH	GC		11	
11q12-q13.1	AE	Serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor)		GS5-10, BPH	GC		11	
11q13	insertion	CCND1: cyclin D1; Primary G1 to S phase cyclins.	not linked with MYC or AR.	Prim, Recur, Mets, vs BPH,	FISH TMA		13	
	insertion	CCND1: cyclin D1; Primary G1 to S phase cyclins.		GS 4-10,BPH	FISH	24.5	15	
	UR	GSTP1 (glutathione S-transferase pi)		GG4/5, BPH,	GC		8	
11q22	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
11q22.3-q23.1	DR	CRYAB (crystallin, a B) heat shock like protein.		GG4/5, BPH,	GC	100	8	
	AE	CRYAB		GS5-10, BPH	GC		11	
	OE	CRYAB	overexpressed in response to ischemia	radical prostatectomy tissue	GC, WB, NB, IH		14	
11q22-q23	DR	ATDC: ataxia-telangiectasia group D-associated protein		GG4/5, BPH,	GC	100	8	
11q 24-q25	AE		Suppression of tumorigenicity 14	GS5-10, BPH,	GC	100	11	
<b>Chromosome 12</b>								
12p11.2	AE	DKFZP56401863 Protein		GS5-10, BPH,	GC	100	11	
12p13.1-p12.3	AE	Microfibril-associated glycoprotein-2		GS5-10, BPH,	GC	100	11	
12p13	DR	KIAK0002/CCND2 (cyclin D2)		GG4/5, BPH,	GC	100	8	
12q	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	22	7	
12q13	DR	KRT5-keratin type II		GG4/5, BPH	GC	100	8	
	UR	KRT18 (Keratin 18)		GS5-9, norm	GC		12	
12q12-q13	UR	Homeo box c8 protein		GG4/5, BPH,	GC	100	8	
	DR	IGFBP6 (insulin-like growth factor binding protein 6)		GG4/5, BPH,	GC	100	8	
	AE	TAR2PT [TAR(HIV) RNA-binding protein 2		GS5-10, BPH	GC		11	
12q13.11-q13.12	DR	NELL2 (nel(chicken)-like2)		GG4/5, BPH,	GC	100	8	
12q13-q14	AE	TS translation elongation factor, mitochondrial		GS5-10, BPH	GC		11	
12q14	AE	RAP1B, member of RAS oncogene family		GS5-10, BPH	GC		11	
12q21	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
12q22-23	AE	Insulin-like growth factor 1 (somatomedia C)		GS5-10, BPH,	GC		11	
	AE	DUS6; dual specificity protein phosphatase 6		GS5-10, BPH	GC		11	
12q24.31	AE	HM74, putative chemokine receptor, GTP-binding protein		GS5-10, BPH	GC		11	
12q24.13	AE	NM23B expressed in NME2		GS5-10, BPH,	GC	100	11	
12q24.4	UR	CAMKK2; calcium/calmodulin-dependant kinase kinase 2 b		GS5-9, BPH	GC		12	
<b>Chromosome 13</b>								
13q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	51	7	
	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	19	7	
	deletion			T1NxMx		6.3	21	
	deletion			T2NO	CGH	com	22	
	LOH	include 68RS20 in Rb gene (13q14)	possible Tumor Suppressor Gene.	Local GS5-7, 8&9, HR, Mets, norm	SB		25	
	LOH	markers selected near Rb1		GS4-9, T2b-T3c, N1,Mx vs norm	PCR	23	33	
	deletion			GS5-10	CGH	55	1	
13q12	deletion	BRCA2 & BRUSH-1	possible Tumor Suppressor Gene.	N/A, 17 malignant tumors, 1 benign	CGH, FISH, PCR		2	
	deletion	Pseudogene of antioncogene nm-23 H1		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
13q14	deletion	Rb(1):Retinoblastoma suppressor gene;	tumor suppressor gene	GS 4-10, BPH	FISH	22	15	
13q14.1-q21.3	deletion		tumor suppressor gene	T1NxMx			21	
13q21	deletion			T2NO	CGH		22	
				T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
				GS5-10	CGH		1	
13q21.1-21.3	deletion			T2NO	CGH		22	
13q22	deletion	Endothelin B receptor gene (ENDRB)		T2NO	CGH	Com.	22	
13q31	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
13q34	AE	Collagen, type IV, alpha1 chain precursor		GS5-10, BPH	GC		11	

**Chromosome 14**

C14	DR	MIG2 (mitogen inducible 2)		GG4/5, BPH,		100	8
14q	deletion			T2NO	CGH		22
14q11.1	AE	Signal transduction protein (SH3 containing)		GS5-10, BPH	GC		11
14q11.2-q12	UR	APEX nuclease		GG4/5, BPH,	GC	100	8
14q11.2-q21	AE	Mitogen-activated protein kinase kinase kinase 5 (MAPAK5)		GS5-10, BPH	GC		11
14q21-q22	DR	PYGL (phosphorylase, glycogen; liver)		GG4/5, BPH,	GC	100	8
14q22-23		PKCh			FISH		4
14q24	DR	TGFB3 (transforming growth factor-b 3		GG4/5, BPH,	GC	100	8
14q31	DR	ITPK1 (Inositol 1,3,4-triphosphate 5/6 kinase)		GG4/5, BPH,	GC	100	8

**Chromosome 15**

15q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	35	7
15q11-q13	UR	Ring Zinc finger protein (ANF127-xp)		GG4/5, BPH,	GC	100	8
15q15-q21.1	DR	FGF7 (fibroblast growth factor 7 [keratinocyte growth factor])		GG4/5, BPH,	GC	100	8
15q15	AE	THBS1, thrombospondin -1		Mets, local, BPH, PIN, norm	GC		5
15q21-q22	DR	LIP2 (lipocortin II)		GG4/5, BPH,	GC	100	8
	DR	ANXA2, annexin 2		GS6-9, BPH, HR	GC		6
15cen-q21	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
	UR	ERK3 (extracellular signal -regulated kinase),at 15q21.		GG4/5, BPH	GC	100	8
15q25-qter	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7
15q26	UR	ALDH6;aldehyde dehydrogenase 1 family member A3		GS5-9, BPH	GC		12

**Chromosome 16**

16p	deletion			N/A, 17 malignant tumor, 1 benign	CGH,FISH,PCR		2
16p11.1	AE	TGFb -1 induced transcript 1		GS5-10, BPH	GC		11
16p11.2	AE	Kinesin-like 4 (replicator binding protein)		GS5-10, BPH,	GC		11
	AE	Protein kinase C, b 1		GS5-10, BPH,	GC		11
16p11-13	AE	NPIP, mRNA, nuclear pore complex interacting protein		GS5-10, BPH,	GC		11
16p12		PKCb1			FISH		4
	AE	G protein coupled receptor, family C, group5, member BPKCb1		GS5-10, BPH,	GC		11
		PKCb1		GS5-10, BPH,	GC		11
16p13.13-p13.12	DR	MYH11(myosin heavy polypeptide 11, smooth muscle)		GG4/5, BPH,	GC	100	8
	DR	A-362G6.1 (hypothetical protein)		GG4/5, BPH	GC	100	8
16q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	46	7
	deletion			pT2NO	CGH		22
	LOH	Using a 16q22-24 highly polymorphic loci	possible tumor suppressor gene	Local, GS5-7, 8&9, HR, Mets, norm	SB	31	25
	deletion			GS5-10	CGH	25	1
	deletion			N/A, 17 malignant tumor, 1 benign	CGH,FISH,PCR		2
16q12	AE	Ste-20 related kinase (to serine threonine kinase 39 at 2q24.3)		GS5-10, BPH	GC		11
16q12.2-q13	DR	KIAA0025 (KIAA025 gene product; MMS inducible gene)		GG4/5, BPH	GC	100	8
16q13	DR	MT1G (metallothionein 1G)		GG4/5, BPH,	GC	100	8
	DR	MT1L (metallothionein 1L)		GG4/5, BPH,	GC	100	8
	DR	MT2A (metallothionein 2A)		GG4/5, BPH,	GC	100	8
	OE	MT1L (metallothionein 1L)	overexpressed in response to ischemia	radical prostatectomy tissue	GC, WB, NB, IH		14
16q13-q21	AE	Matrix metalloproteinase 2		GS5-10, BPH,	GC		11
16q22	AE	Dihydroorotate dehydrogenase		GS5-10, BPH	GC		11
	deletion			GS5-10	CGH		1
16q22.1	deletion	CDH1 (E-cadherin ) or CTCF gene (regulates MYC expression)	possible tumor suppressor gene	T2NO	CGH		22
	DR	CDH1 (E-cadherin )		prostate cancer	GC		5
16q23.1	AE	Carboxylesterase 2 (intestine, liver)		GS5-10, BPH	GC		11
16q23-q24	LOH		possible tumor suppressor gene	T3NoMo	PCR MSA	53	16
16q24	deletion		possible tumor suppressor gene	T1-4,Nx,Mx,0,1 v norm	CGH& FISH		7

**Chromosome 17**

C17	insertion insertion insertion UR	Polysomy Her-2/neu oncogenes .		T3, GS7 GP4, GS7, <T3b	ISH FISH	Rare Com	58 54
17p	deletion insertion insertion deletion	PYCR1/pyrroline 5-carboxylate reductase 1	aneusomy in metastases	GG4/5, BPH, T1-4,Nx,Mx,0,1 vs norm T1NxMx T2NO	GC CGH & FISH CGH	100 41 10.4	8 7 21
17p12	insertion		Possible oncogenes	N/A, 17 malignant tumors, 1 benign T1NxMx	CGH,FISH,PCR CGH		2 21
17p13	deletion deletion	p53 p53:	50% loss in AR and 65% in HR	T(2-3) N1(-3)M(0) GS4-9, Lymph Node Mets. & Mets, HR (GS6-10) vs TURP& blood	FISH CGH, SB & PCR MSA	46 50/65	56 3
17p13.1	LOH LOH LOH DR	p53 p53 p53; deactivation of the gene SERPINF1 (serine [or cysteine] proteinase inhibitor, clade F, 1)	possible anti-metastasis gene	Prim vs Mets (GS5-9)(not BPH or norm). norm. GS 3/3 to 4/5	IH PCR/seq for SNPs. IH, PCR	8.5 vs23' >50	34 35 36
17q	insertion insertion deletion			GG4/5, BPH, T1NxMx T2NO	GC CGH CGH	100 10.4	8 21 22
17q11.2	insertion		Possible oncogene	N/A, 17 malignant tumors, 1 benign GS5-10	CGH,FISH,PCR CGH		2 1
17q12-q21	DR	KRT14 (keratin14)		GG4/5, BPH, GS 4-10,BPH	GC FISH	100	8 15
17q13	LOH AE	HER-2/neu c-ERBB2; TNF (ligand) superfamily, member 12	disputed cf ref 13	GS5-10, BPH, GS5-10, BPH, GS5-10	GC GC CGH		11 11 1
17q	AE insertion	COP22 for nonclathrin coat protein zeta-COP		GS5-10, BPH T2NO	GC CGH	23	11 22
17q21	AE	Amine oxidase, (vascular adhesion protein 1)		GS5-10, BPH	GC		11
17q21.1	insertion	ERBB2 (breast)?	Possible oncogene	T2NO T1-4,Nx,Mx,0,1 vs norm	CGH CGH & FISH		22 7
17q21.3	AE AE	NM23A expressed in NME CDC6 (cell division cycle 6, S. cerevisiae) homolog		Prim, Recur, Mets, vs BPH GS5-10, BPH,	FISH TMA GC		13 11
17q24	LOH	PKCa:?			FISH		4
17q24-qter	insertion insertion		Possible oncogene oncogene	T1NxMx GS5-10 GS5-9	CGH CGH GC, IH		21 1 12
17q25	UR UR UR	FASN;fatty acid synthase P4HB;procollagen-proline 2-oxoglutarate 4 dioxygenase beta FASN;fatty acid synthase		GS5-9 Mets, local, BPH, PIN, norm	GC GC		12 12 5
17q25.3	AE UR	TIMP2, tissue inhibitor of metalloproteinase 2 PCYR1; pyrroline-5carboxylate reductase 1		Mets, local, BPH, PIN, norm GS5-9, normal	GC GC		5 12

Chromosome	Region	Variant	Gene/Function	Associated Genes	Method	Value 1	Value 2
<b>Chromosome 18</b>							
C18 long		deletion	DCC and DPC4 (Smad 4);	2 possible tumor suppressor genes	T2NO	CGH	22
18q		deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	19 7
		insertion			T1-4,Nx,Mx,0,1 vs norm		30
		deletion			T2NO	CGH	22
		deletion			T1NxMx	CGH	4.2 21
		deletion			GS5-10	CGH	20 1
18q11-q12		DR	SLC14A1-urea transporter		GG4/5, BPH,	GC	100 8
18q12		insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	7
18q21		deletion	DCC, DPC4 or ?	possible tumor suppressor genes	GS5-10	CGH	1
18q21-22		LOH			T3NoMo	PCR MSA	45 16
18q21.3		UR	Bcl-2		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	7
		deletion	DCC	tumorigenesis			
			SMAD4 (DPC4)				10./13 37, 38
		UR	Bcl-2		HR vs non-HR mets	IH	24
		UR	P15- protease inhibitor 5 (maspin)		GG4/5, BPH,	GC	100 8
18q22-qter		deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	7
<b>Chromosome 19</b>							
C19		deletion	Contains Putative TSG's; BAX & LKB1		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	43 7
19p13.1-13.2		UR	MIC-1;PLAB		GG4/5, BPH	GC	100 8
		UR	MIC-1;PLAB;secreted cytokine		GS5-9, norm	GC, RTPCR	12
19p13.2		AE	CipP (caseinolytic protease, ATP-dependant) homolog		GS5-10, BPH	GC	11
19p13.3			HIC-1; hypermethylated in cancer	possible tumor suppressor gene			26, 39
19q11-q13.2		UR	Hepsin: transmembrane extracellular serine protease;	cell growth control	NBPM;	GC & ISH	40
		UR	Hepsin		GG4/5, BPH	GC	100 8
		UR	Hepsin		GS5-10, BPH	GC, RTPCR	100 11
		UR	Hepsin		GS5-9, norm	GC, RTPCR,	12
			hepsin (HPN)		prostate cancer tissue, BPH	GC	5
19q13.1		DR	COX7A1 (cytochrome c oxidase subunit V11a polypeptide 1)		GG4/5, BPH,	GC	100 8
		AE	FXYD domain-containing ion transporter regulator 1		GS5-10, BPH,	GC	11
19pter-q13.1		deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	7
19q13.3		AE	Fc fragment of IgG, receptor, transporter, alpha		GS5-10, BPH	GC	11
19q13.3-qter		DR	RRAS (related RAS viral [r-ras] oncogene homology		GG4/5, BPH	GC	100 8
19q13.4		LOH	PKCg ;			FISH	4
		UR	MTA-1, alpha-1-B glycoprotein		Mets, local, BPH, PIN, norm	GC	5
19q13.41			PSA (KLK3); serine protease				41
19q13.3-q13.4			KLK1;pancreatic/renal kalikrein				41
			KLK2 human glandular kalikrein				41
			PRSSL1(NES1);normal epithelial cell-specific 1	breast.			43, 44, 42
			Protease M (zyme, neurosin)	brain, ovarian and breast cancer			43, 45, 46, 47, 42
			KLK4; prostate specific antigen-related serine protease gene,	endometrial, prostate et al	N/A	PCR	42
		DR	KLK5		norm, T1-T3	RT-PCR	48
		AE	KLK14; secreted serine protease with trypsin-like substrate.	follow klk3 and klk2	benign, malignant	ISH	49
			TLSP; trypsin-like serine protease				41
			HSCCE; human stratum corneum chymotryptic enzyme.				41
			KLK-L1-4				41

<b>Chromosome 20</b>								
20pter-p12	DR	PRNP (prion protein (p27-30))		GG4/5, BPH,	GC	100	8	
	AE	Chromogranin B (secretogranin 1)		GS5-10, BPH,	GC		11	
20p11.2	AE	Transglutaminase 3		GS5-10, BPH,	GC		11	
20q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	22	7	
	insertion			T2NO	CGH		22	
	deletion			N/A, 17 malignant tumor, 1 benign	CGH, FISH, PCR		2	
20cen-q22	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
20q11.2	UR	FLS353; Chromosome 20 open reading frame 1; unknown		Mets, local, BPH, PIN, norm	GC		5	
20q12-q13	AE	Phospholipid transfer protein (PLTP)		GS5-10, BPH	GC		11	
20q13.1	UR	MYBL2, v-myb myeloblastosis viral oncogene		Mets, local, BPH, PIN, norm	GC		5	
20q13.11	AE	Ubiquitin carrier protein EC-2		GS5-10, BPH,	GC		11	
20q13.2-q13.3	insertion	CYP24 , ZNF217 and NABC1; (in breast cancer)	possible oncogene	T2NO	CGH		22	
<b>Chromosome 21</b>								
21q22	AE	Collagen, type VI, alpha 1H2BFS;		GS5-10, BPH,	GC		11	
	UR	H2B Histone, family member 5		GS5-9, normal	GC		12	
21q22.13	UR	SIM2; single-minded homolog 2 (Drosophila)		GS5-9, BPH	GC		12	
21q22.13-22.2	AE	KCNJ6 (potassium inwardly rectifying channel, J, mem 6).		GS5-10, BPH	GC		11	
<b>Chromosome 22</b>								
22q	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	46	7	
22q12.3	DR	TIMP3(tissue inhibitor of metalloproteinase 3		GG4/5, BPH,	GC	100	8	
22q13	deletion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
	AE	Thiosulfate sulfurtransferase (rhodanese)		GS5-10, BPH	GC		11	
22q13.31	UR	NBK apoptotic inducer protein		GG4/5, BPH	GC	100	8	
<b>X Chromosome</b>								
Xp	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	24	7	
	insertion			GS5-10	CGH	16	1	
Xp11.23	DR	PLP2 (proteolipid protein 2[colonic epithelium enriched])		GG4/5, BPH,	GC	100	8	
Xpter-xp21	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH		7	
Xp21.1	DR	SRPX (sushi-repeat-containing protein, X chromosome).		GG4/5, BPH,	GC	100	8	
	AE	Cytochrome b-245, beta polypeptide		GS5-10, BPH	GC		11	
Xp21.13	AE	Thioredoxin peroxidase (antioxidant enzyme)		GS5-10, BPH,	GC		11	
Xp22.2	DR	GPM6B (glycoprotein M6B)		GG4/5, BPH,	GC	100	8	
	AE	GPM6B (glycoprotein M6B)		GS5-10, BPH,	GC		11	
Xp22.3	AE	Transducin (beta)-lie 1		GS5-10, BPH,	GC		11	
Xp22.32	UR	SLC25A6/solute carrier family 25, member A6		GG4/5, BPH,	GC	100	8	
Xq	insertion			T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	35	7	
	insertion			GS5-10	CGH	23	1	
Xq12	insertion	Androgen Receptor ;		T1-4,Nx,Mx,0,1 vs norm	CGH& FISH,		7	
	AE	nuclear transcription factorEphrin-B1 (EFNB1)			IH	50, 51		
	insertion	Androgen Receptor ;	metastases & hormone refractory	Prim, Recur, Mets, vs BPH	FISH TMA	20	13	
	LOH	Androgen Receptor ;		Prim, Mets (D2), norm, cell line	PCR		52	
Xq12-q13	insertion		possible oncogene	GS5-10	CGH		1	
	insertion		Found in metastses only	GS5-10	CGH	2 /.15	1	
Xq13.1	UR	GJB1/gap junction protein		GG4/5, BPH,	GC	100	8	
Xcen-q13	insertion		Hormone refractory tumors only	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	100	7	
Xq23-qter	deletion		Hormone refractory tumors only	T1-4,Nx,Mx,0,1 vs norm	CGH& FISH	100	7	
Xq21.3		Pseudogene of PKCi?			FISH		4	
Xq21.33-q22.3	UR	TMSNB/NB thymosin b.		GG4/5, BPH,	GC	100	8	
Xq26-q25	AE	COVA 1(cytosolic ovarian carcinoma antigen 1)		GS5-10, BPH,	GC	100	11	
Xq28	AE	Glucose-6-phosphate dehydrogenase G6PD		GS5-10, BPH,	GC		11	
<b>Y Chromosome</b>								
Y	deletion			GS5-10	CGH	16	1	
	deletion			N/A, 17malinant tumor, 1 benign	CGH,FISH,PCR		2	
Yp	UR	SLC25A6/solute carrier family 25, member A6		GG4/5, BPH,	GC	100	8	