

# The effects of short-term genistein intervention on prostate biomarker expression in patients with localised prostate cancer before radical prostatectomy

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#### **Abstract**

Nutritionally relevant levels of genistein, the predominant isoflavone in soyabean associated with lower risk of prostate cancer (PCa), may modulate the expression of prostate tissue biomarkers associated with cancer prediction and progression. A phase 2 placebocontrolled, randomised, double-blind clinical trial was conducted in forty-seven Norwegian patients before prostatectomy. Intervention was 30 mg genistein or placebo capsules daily for 3-6 weeks. Luminal cells from malignant and benign glands were isolated with laser capture microdissection and the mRNA levels of androgen-related biomarkers (androgen receptor, NK3 homeobox 1, kallikrein-related peptide 4 (KLK4)) and cell cycle-related genes ( $p21^{Waf1/Cip1}$ ,  $p27^{Kip1}$ , p53) were analysed with real-time semiquantitative PCR. Immunohistochemistry of androgen-, cell cycle-, proliferative- (Ki67 nuclear antigen), apoptotic- (B-cell CLL/lymphoma 2 (BCL-2) and BCL-2-associated X protein) and neuroendocrine differentiation-related biomarkers (neuron-specific enolase and cytoplasmic chromogranin A) was performed using tissue microarrays containing normal, Gleason grade 3 and grade 4 prostate tissues. There were no significant effects by genistein intervention on proliferation-, cell cycle-, apoptosis- or neuroendocrine biomarkers. Genistein intervention, however, significantly reduced the mRNA level of KLK4 in tumour cells (P=0·033) and there was a nonsignificant reduction in androgen and cell cycle-related biomarkers, except for p27Kip1, whose expression in the nuclear compartment was increased. Genistein intervention modulated the expression of several biomarkers which may be related to PCa prediction and progression. The present study supports genistein as a chemopreventive agent in PCa. Further investigation is warranted in larger and longer-duration studies.

Key words: Biomarkers: Genistein: Immunohistochemistry: PCR: Prostate cancer

Although not an ideal biomarker, serum prostate specific antigen is the only predictive and prognostic prostate cancer (PCa) biomarker widely used in clinical practice. Several proteins have been proposed as candidate biomarkers exhibiting all or some of these predictive or prognostic properties (1-3). Among these are androgen-, cell cycle-, proliferation-, apoptosis- and neuroendocrine differentiation-related biomarkers.

The soya isoflavone, genistein, is a promising chemopreventive agent in PCa based on molecular, epidemiological and clinical studies (4,5). We have previously reported the clinical endpoints of a phase 2 clinical randomised trial with short-term genistein intervention in patients with localised PCa. Genistein reduced the level of serum prostate specific antigen without any effects on hormones<sup>(6)</sup>. The purpose of

Abbreviations: AR, androgen receptor; BAX, B-cell CLL/lymphoma 2-associated X protein; BCL-2, B-cell CLL/lymphoma 2; CgA, cytoplasmic chromogranin A;  $G3, Gleason\ grade\ 4;\ KLK4,\ kallikrein-related\ peptide\ 4;\ NKX3\cdot 1,\ NK3\ homeobox\ 1;\ NSE,\ neuron-specific\ enolase;\ p21^{Waf1/Cip1},\ cyclin-particle particle par$ dependent kinase inhibitor 1A; p27<sup>Kip1</sup>, cyclin-dependent kinase inhibitor 1B; p53, tumour protein p53; PCa, prostate cancer.



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Table 1. Primers for PCR

Gene	Forward primer	Reverse primer	
AR KLK4 NKX3-1 p21 p27 p53 ALAS1	TGGAAGCTGCAAGGTCTTCT ATGGAAAACGAATTGTTCTGCTC GGCCTGGGAGTCTCTTGACTCCACTAC AGGGGACAGCAGAGGAAGA TCAGAAGACGTCAAACGTAAACA CCCCAGCCAAAGAAGAAAC CTGCAAAGATCTGACCCCTC	AAGCCTCTCCTTCCTCTGT CAGCACGGTAGGCATTCTG ATGTGGAGCCCAAACCACAGAAAAATG CTTCCTGTGGGCGGATTAG CCAACGCTTTTAGAGGCAGA TTTTTATGGCGGGAGGTAGA CCTCATCCACGAAGGTGATT	

AR, androgen receptor; KLK4, kallikrein-related peptide 4; NKX3·1, NK3 homeobox 1; p21, cyclin-dependent kinase inhibitor 1A; p27, cyclin-dependent kinase inhibitor 1B; p53, tumour protein p53; ALASI, aminolevulinate synthase gene.

the present study was to investigate the modulation of potential candidate PCa tissue biomarkers by genistein at a dose that can be obtained from a diet rich in soya-based food.

# **Experimental methods**

# Patients and study design

A total of forty-seven patients with localised PCa scheduled to be treated by radical prostatectomy were randomised during April 2007-August 2008. Next, a total of forty-one prostates were analysed and forty were considered as evaluable according to the study protocol. However, one specimen only contained enough tumour material for pathological analysis and could therefore not be analysed for biomarkers. The study was a single-centre, placebo-controlled, randomised and double-blind phase 2 clinical trial with two treatment arms<sup>(6)</sup>. One arm was given 30 mg synthetic genistein (geniVida®(7)) daily and one arm received placebo. The intervention period was 3-6 weeks (average 33d) before radical prostatectomy. This study was conducted according to the guidelines laid down in the Declaration of Helsinki and all procedures involving human subjects/patients were approved by the Norwegian Medicines Agency, the Regional Ethics Committee, the Privacy Ombudsman and the Prostate Biobank at the Oslo University Hospital, Aker. Written informed consent was obtained from all subjects/patients. The study has been registered in the ClinicalTrials.gov registry (study identifier: NCT00546039).



After surgery, the prostates were immediately transferred to the pathology department and macro-dissected by a pathologist. Fresh tissue cubes, approximately  $0.5\,\mathrm{cm}^3$ , were cut from tumour and normal tissue and frozen at  $-80^\circ\mathrm{C}$ . Frozen  $10\,\mu\mathrm{m}$  sections were stained with haematoxylin and eosin. At least  $10\,000$  cells from histologically benign and cancer glands from each patient were selected and captured using laser capture microdissection (Leica LMD 6000; Leica Microsystems) and frozen at  $-80^\circ\mathrm{C}^{(8)}$ .

Total RNA was isolated using the Arcturus PicoPure RNA Isolation Kit (Molecular Devices) according to the manufacturer's protocol. The concentration of RNA was determined by measurements in the NanoDrop ND-1000 spectrophotometer (NanoDrop Technologies). Reverse transcription of

50 ng total RNA was performed with the iScript cDNA Synthesis Kit (Bio-Rad) according to the manufacturer's protocol. Although fresh prostate tissue was acquired from most patients, tumour was found in only ten study subjects from the genistein arm and twelve from the placebo arm.

#### Semi-quantitative real-time RT PCR

Primers were designed with Primer3 provided by the Whitehead Institute for Biomedical Research (http://fokker.wi.mit.edu/cgibin/primer3/primer3\_www.cgi). Sequences are listed in Table 1. Relative quantification of gene expression by semi-quantitative real-time RT-PCR was performed by analysing the expression of human housekeeping delta-aminolevulinate synthase gene. Real-time PCR was performed on the Bio-Rad Opticon DNA Engine. All amplifications were run as triplicates with standard curves. Here, 1 ng of template was used in each reaction following the manufacturer's protocol (Bio-Rad SYBR Green Master Mix). The same set-up was used for all primers (sixty-six cycles of 55°C annealing, 72°C extension and 90°C denaturing).

#### Tissue processing and immunohistochemical analysis

The prostates were fixed in 10% buffered formalin, macrodissected and paraffin-embedded. Tissue microarrays were

Table 2. Antibodies for immunohistochemistry

Antigen	Clone	Supplier	Dilution	Treatment
AR KLK4 NKX3·1 p21 p27 p53 Ki67 BAX BCL-2 NSE CgA	MS-443 Rabbit Rabbit EA-10 SC-528 Bp53-11 30-9 2D2 124 E27 LK2H10	Thermo Scientific F.S lab F.S lab Calbiochem Santa Cruz Roche Roche Thermo Scientific Roche Roche Roche Roche	Prediluted 3 µg/ml 1 µg/ml 0.25 µg/ml 1 µg/ml Prediluted Prediluted 12.5 µg/ml Prediluted Prediluted Prediluted	Benchmark XT Autoclave Autoclave PT-module* PT-Module† Benchmark XT Benchmark XT PT-module‡ Benchmark XT Benchmark XT Benchmark XT

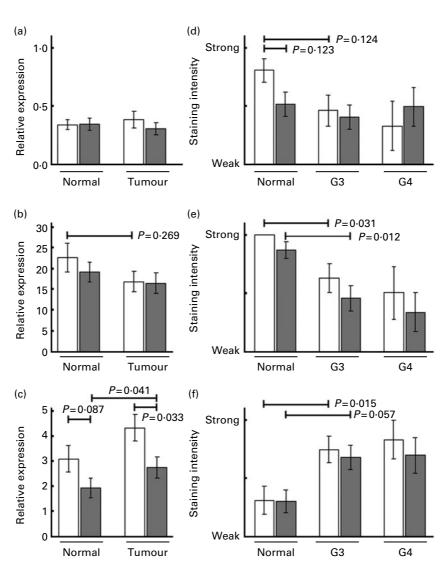
AR, androgen receptor; KLK4, kallikrein-related peptide 4; NKX3·1, NK3 homeobox 1; p21, cyclin-dependent kinase inhibitor 1A; p27, cyclin-dependent kinase inhibitor 1B; p53, tumour protein p53; BAX, B-cell CLL/lymphoma 2-associated X protein; BCL-2, B-cell CLL/lymphoma 2; NSE, neuron-specific enolase; CgA, cytoplasmic chromogranin A.

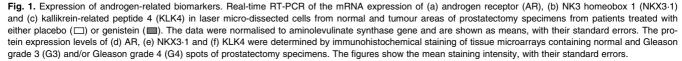


<sup>\*</sup> Citrate buffer pH 6.

<sup>†</sup>Tris buffer pH 9.

<sup>‡</sup>EDTA buffer pH 8.





assembled using the TMABooster (Alphelys) sampler. Normal, Gleason grade 3 (G3) and Gleason grade 4 (G4), if available, prostate tumours were marked on haematoxylin/ eosin-stained sections. From each patient, three 0.6 mm biopsy cores were taken from normal prostatic tissue and each available Gleason grade tumour. The sampling size of three cores has previously been shown to be representative of PCa<sup>(9)</sup>. Next, 10 µm tissue microarray sections on glass slides (SuperFrost®) were treated according to the procedures listed in Table 2. 'Benchmark XT' used automated immunohistochemical staining (Benchmark XT; Ventana Medical Systems). The slides were treated in a dry cabinet for 60 min at 60°C, cooled and washed with a detergent before conditioning with cell conditioning 1 (CC1) buffer (Ventana Medical Systems). Detection of the bound primary antibody was visualised with the iView™ DAB Detection kit

(Ventana Medical Systems). The cells were counterstained with haematoxylin and Bluing reagent (Ventana Medical Systems) for 4 min. 'Autoclave' used citrate buffer pH 6.4 at 120°C for 40 min for antigen retrieval. Immunohistochemistry was performed according to the manufacturer's protocol (BioGenex Detection kit). The slides were visualised by diaminobenzidine staining followed by haematoxylin staining. 'PT-module' used different buffers according to Table 2. The slides were treated in 98°C for 25 min in the PT-Module (Lab Vision Corporation) for antigen retrieval. Immunohistochemistry and visualisation were performed according to the manufacturer's protocol (UltraVision ONE Detection System; Thermo Fisher Scientific). The cells were counterstained with haematoxylin. The stained tissues were then independently analysed by two consultant pathologists (A. S. and C. H.), who later made a consensus evaluation



on each sample. Samples not containing the designated Gleason grade were not analysed and treated as missing.

The staining intensities of nuclear androgen receptor (AR), cytoplasmic and nuclear NK3 homeobox 1 (NKX3·1), cytoplasmic kallikrein-related peptide 4 (KLK4), cytoplasmic and nuclear cyclin-dependent kinase inhibitor 1B (p27Kip1) and cytoplasmic B-cell CLL/lymphoma 2-associated X protein (BAX) were scored as either weak/moderate (0) or strong (1), whereas the staining intensities of cytoplasmic B-cell CLL/ lymphoma 2 (BCL-2), cytoplasmic neuron-specific enolase (NSE: also called ENO2) and cytoplasmic chromogranin A (CgA) were scored as negative (0) or positive cytoplasmic staining (1).

Sample sizes for the placebo arm were: sixteen containing normal glandular tissue, sixteen G3 and six G4 tumours; and for the genistein arm: twenty-three normal glandular tissue, twenty-two G3 and ten G4. In the placebo arm, one G3 disappeared due to technical reasons for AR and p27<sup>Kip1</sup>. In the genistein arm, one G4 disappeared for NKX3·1 and cyclin-dependent kinase inhibitor 1A (p $21^{Waf1/Cip1}$ ).

# Statistical analysis

Values are expressed as means with their standard errors for continuous and ordinal data. Differences between two related and independent samples were tested by Fisher-Pitman permutation tests for paired and two-sample interval-scaled data. The P-values were based on 2000 simulations and considered significant at P < 0.050. The analyses were carried out in STATA/IC 11.1 for Windows (32-bit) and Sigmaplot 11.0 for Windows was used to create the figures.

#### **Results**

#### Androgen-related biomarkers

Genistein intervention significantly reduced KLK4 mRNA expression in tumour cells (P=0.033). The down-regulation of AR protein expression and KLK4 mRNA level in normal cells were not statistically significant (P=0.123 and P=0.087; Fig. 1(c) and (d)). There was a general non-significant tendency by genistein intervention to reduce the expression of androgen-related biomarkers (Fig. 1).

The AR and NKX3·1 nuclear protein expression in both study arms were reduced in higher Gleason grades, whereas both the mRNA and protein expression were increased for KLK4. AR mRNA expression was unchanged. Fig. 2(a) depicts a patient treated with placebo showing strong nuclear AR staining intensity in normal cells and weak intensity in the case of tumour. The cytoplasmic expression of NKX3·1 was equivalently increased (data not shown). Fig. 2(b) shows a patient from the genistein group with strong nuclear/weak cytoplasmic NKX3·1 staining intensity in normal cells and weak nuclear/strong cytoplasmic staining intensity in tumour. Fig. 2(c) depicts a patient treated with genistein showing weak cytoplasmic KLK4 staining intensity in normal cells and strong intensity in tumour.

# Cell cycle-related biomarkers

Genistein intervention had no significant effects on p21 Waf1/Cip1 p27<sup>Kip1</sup> or tumour protein p53 (p53) mRNA and protein expression (Fig. 3). There was a non-significant reduction in

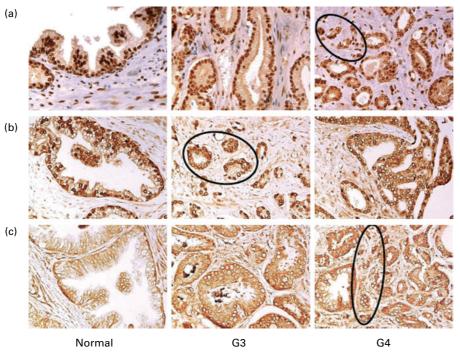
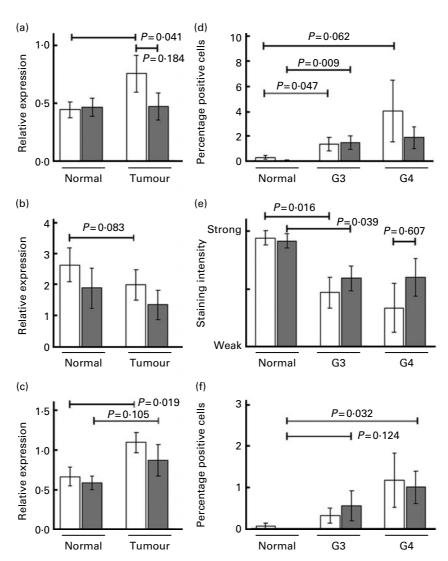
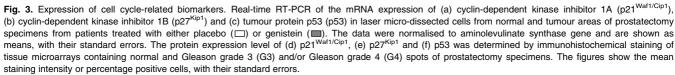


Fig. 2. Immunohistochemical staining of androgen-related biomarkers. Each row shows tissue microarray spots from a single patient containing normal, Gleason grade 3 (G3) and Gleason grade 4 (G4) prostate tissues. Encircled areas contain designated Gleason grade. Magnification ×400. (a) Androgen receptor, (b) NK3 homeobox 1 and (c) kallikrein-related peptide 4.



2142 B. Lazarevic et al.





the expression of p21<sup>Waf1/Cip1</sup> mRNA expression in tumour (P=0·184) and a slight reduction in p27<sup>Kip1</sup> and p53 mRNA expression, whereas 27<sup>kip1</sup> protein nuclear expression was slightly increased in G3 and G4 in the genistein arm compared with placebo.

The mRNA and protein expression results for the cell cycle-related biomarkers were coherent in both study arms, showing increased levels of p21<sup>Waf1/Cip1</sup> and p53 and reduced levels of p27<sup>Kip1</sup> in tumour tissue compared to normal tissue. The percentage p21<sup>Waf1/Cip1</sup> positive cells was less than 1% in normal prostate tissue and increased with increasing Gleason grade to about 2% in G3 cells and 4% in G4 cells. Fig. 4(a) depicts a patient from the genistein group showing an increasing number of p21<sup>Waf1/Cip1</sup> positive nuclei from normal tissue to G4. The nuclear expression of p27<sup>Kip1</sup> was significantly reduced in G3 (*P*=0·016) compared to normal, whereas the

mRNA level was non-significantly reduced in tumour compared to normal cells (Fig. 3(b) and (e)). Fig. 4(b) shows a patient from the placebo group with strong nuclear/weak cytoplasmic staining intensity in normal cells and weak nuclear/strong cytoplasmic p27<sup>Kip1</sup> staining in G3. Fig. 4(c) shows a patient treated with placebo with no p53 positive cells in normal tissue and with single p53 positive cells in G3 and G4 tissues.

#### Proliferation- and apoptosis-related biomarkers

Genistein intervention had no significant effects on the protein expression of Ki67, BAX or BCL-2 (Fig. 5).

In both study arms, Ki67 and BAX expression increased with increasing Gleason grade. Ki67 was expressed by 1% of normal epithelial cells and it increased significantly to 3%



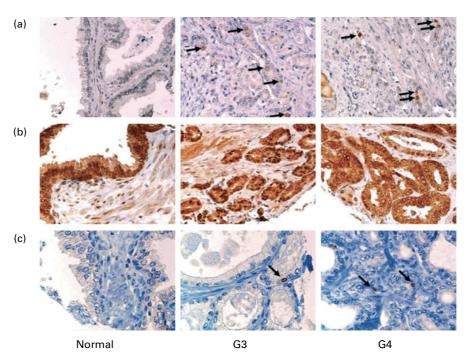


Fig. 4. Immunohistochemical staining of cell cycle-related biomarkers. Each row shows tissue microarray spots from a single patient containing normal, Gleason grade 3 (G3) and/or Gleason grade 4 (G4) prostate tissues. Magnification ×400 for (a) cyclin-dependent kinase inhibitor 1A and (b) cyclin-dependent kinase inhibitor 1B. Magnification ×600 for (c) tumour protein p53.

in G3 cells (P < 0.001) and further to approximately 5% in G4 cells (Fig. 5(a)). Fig. 6(a) depicts a patient treated with genistein showing increasing number of Ki67 positive cells from normal tissue to G4. BAX protein expression increased significantly (P=0.011) in G3 compared to normal cells (Fig. 5(b)). Fig. 6(b) shows a patient treated with genistein with weak cytoplasmic BAX staining in normal tissue and increasing intensity in tumour tissue. BCL-2 was in general not expressed in normal epithelial cytoplasm (Fig. 5(c)). The increased expression in malignant tissue was not statistically

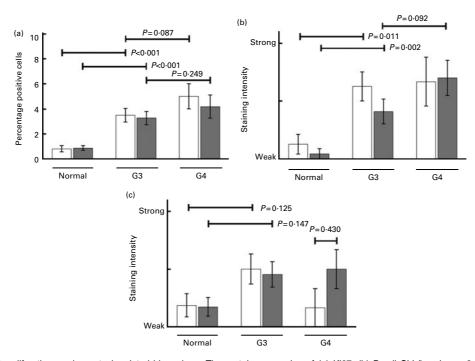


Fig. 5. Expression of proliferation- and apoptosis-related biomarkers. The protein expression of (a) Ki67, (b) B-cell CLL/lymphoma 2-associated X protein and (c) B-cell CLL/lymphoma 2 was determined by immunohistochemical staining of tissue microarrays containing normal and Gleason grade 3 (G3) and/or Gleason grade 4 (G4) spots of prostatectomy specimens. The figures show the mean staining intensity or percentage positive cells, with their standard errors.  $\Box$ , Placebo; , genistein.



2144

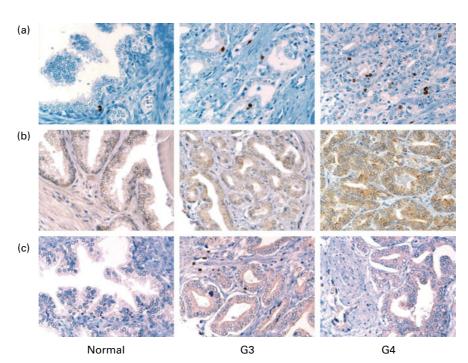


Fig. 6. Immunohistochemical staining of proliferation- and apoptosis-related biomarkers. Each row shows tissue microarray spots from a single patient containing normal, Gleason grade 3 (G3) and Gleason grade 4 (G4) prostate tissues. Magnification ×400. (a) Ki67, (b) B-cell CLL/lymphoma 2-associated X protein and (c) B-cell CLL/lymphoma.

significant (P=0·125). Fig. 6(b) shows a patient treated with genistein with weak BCL-2 staining in normal tissue and stronger staining in tumour tissue.

## Neuroendocrine differentiation-related biomarkers

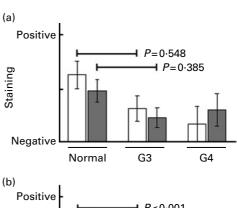
Genistein intervention had no significant effect on NSE or CgA (Fig. 7).

In both study arms, the expression of the neuroendocrine differentiation-related biomarkers indicated reduced levels with increasing Gleason grade. Fig. 8(a) shows a patient treated with placebo with decreasing number of NSE-positive cells in G3 tumour. There was a clear presence of CgA-positive cells in normal tissue, which was completely abolished in G4 tissue in both treatment arms (P<0.001; Fig. 7(b)). Fig. 8(b) shows a patient treated with genistein with decreasing number of CgA-positive cells with higher Gleason grade.

## Discussion

To our knowledge, this is the first study which exclusively investigates the effects by genistein alone in men with PCa. We showed that genistein intervention significantly downregulated the mRNA expression levels of KLK4 in cancer cells in men with localised PCa. Genistein has previously been shown to decrease AR nuclear binding to the transcriptional binding site for AR<sup>(10)</sup>. We also report a decrease by genistein on most of the selected androgen-related biomarkers, corroborating the results of our *in vitro* studies<sup>(11)</sup>. However, the average plasma level in the present study was at least 25-fold lower than the concentrations used in cell culture studies, indicating that in vivo nutritional relevant levels

of genistein may have comparable effects to high-dose cell studies. This may be related to longer in vivo exposure. The androgen-related biomarkers used in this study have been attributed roles in the development or progression of PCa. NKX3·1 is an androgen-regulated homeobox gene located



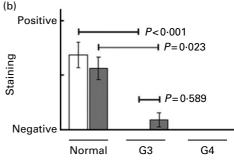


Fig. 7. Expression of neuroendocrine differentiation-related biomarkers. The protein expression of (a) neuron-specific enolase and (b) cytoplasmic chromogranin A was determined by immunohistochemical staining of tissue microarrays containing normal and Gleason grade 3 (G3) and/or Gleason grade 4 (G4) spots of prostatectomy specimens. The figures show the mean positive/ negative stain, with their standard errors. 

, Placebo; 
, genistein.





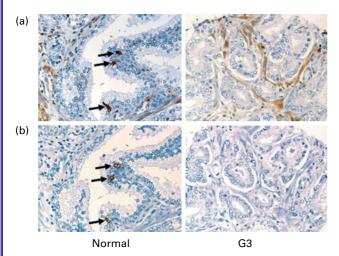


Fig. 8. Immunohistochemical staining of neuroendocrine differentiationrelated biomarkers. Both rows show tissue microarray spots from the same patient containing normal and Gleason grade 3 (G3) prostate tissues. Magnification  $\times 400$ . (a) neuron-specific enolase and (b) cytoplasmic chromogranin A.

on chromosome 8p21·2, a region that shows a high degree of loss of heterozygosity in PCa<sup>(12,13)</sup>. Reports of its function and expression in PCa have been conflicting. However, most reports indicate that NKX3·1 acts as a non-classical tumour suppressor and that its expression is reduced in primary PCa and further reduced in metastatic PCa<sup>(14,15)</sup>. However, our immunohistochemistry results indicated a translocation of NKX3·1 from the nucleus to the cytoplasm and not a total reduction with increasing Gleason grade. Interestingly, increased cytoplasmic stain intensity in primary PCa was briefly mentioned by Gurel et al. (15), although it was not taken into account during scoring. Different splicing variants of NKX3·1 and antibodies have previously been attributed for the conflicting results of NKX3·1 expression. Human kallikrein 4 (KLK4), a serine protease belonging to the prostate specific antigen-related kallikrein family, is overexpressed in PCa and might be a proliferative factor acting directly or indirectly on cell cycle regulators (16). Increased KLK4 levels in PC-3 cells have also been associated with a transcriptional repression of E-cadherin and increased Matrigel motility<sup>(17)</sup>. Our results were consistent with previous reports showing an overexpression of KLK4 mRNA and protein in PCa.

There are no previous reports on in vivo studies on the effects of genistein on the cell cycle-related genes p21 Waf1/Cip1, p27 Kip1 and p53. However, in vitro studies on prostate, breast and lung cancer cells report that genistein up-regulates them, in line with its inhibitory effect on cell cycle progression (18,19). We did not observe any significant effects by genistein, but the mRNA levels of p21 Waf1/Cip1, p27Kip1 and p53 were slightly down-regulated. The reason for this discrepancy is not clear. Interestingly, we showed an increased expression of p21Waf1/Cip1 and p53 mRNA and protein expression in G3 and that the expression may further increase with increasing Gleason grade. Navone et al. (20) showed that p53 accumulation is associated with late-stage PCa. With a cutoff value at 5% of p53 positive cells, they detected no accumulation in nineteen patients with Gleason score 5-7, whereas nineteen of forty-two patients with Gleason score 8-10 showed accumulation. p21Waf1/Cip1 may act both as a cell cycle negative regulator and anti-apoptotic mediator and its protein expression has been shown to correlate strongly with p53 in PCa<sup>(21,22)</sup>. The reduction of p27<sup>Kip1</sup> mRNA and nuclear protein expression in both study arms and the small reversal of protein expression in G3 and G4 by genistein intervention may have an interesting implication for the cell cycle modulating properties of genistein. Loss of phosphatase and tensin homologue function and/or constitutive activation of Akt/protein kinase B by the phosphoinositide kinase-3 pathway, commonly found in PCa, will reduce the expression of p27<sup>Kip1 (23,24)</sup>. In vitro studies on genistein indicate that it inhibits protein kinase B constitutive activation by re-establishing phosphatase and tensin homologue expression<sup>(25,26)</sup>. Although the increase of p27<sup>kip1</sup> in G3 and G4 by genistein was not significant, this result may indicate the mechanism for its G<sub>1</sub> cell cycle arrest and proliferation inhibition of tumour cells<sup>(27)</sup>. Further research is needed in this connection.

The proliferation biomarker Ki67 was clearly up-regulated in tumour tissue, although it was not modulated by genistein. Our detection of 1% Ki67 positive luminal cells in normal and 4-6% in tumour tissue corroborates with previous publications<sup>(28)</sup>. The increasing levels of Ki67, p21<sup>Waf1/Cip1</sup> and p53 may indicate that loss of p53 function already is present in some G3 cancers.

The apoptotic biomarkers BAX and BCL-2 were both significantly or near-significantly up-regulated in PCa, corroborating a previous report on Gleason score  $5-10^{(29)}$ . We observed no significant regulation of BAX or BCL-2 by genistein intervention.

The role of neuroendocrine markers as prognostic factors is controversial, although the serum level of CgA and NSE seems to have a prognostic value in castration-resistant PCa<sup>(30-32)</sup>. Genistein treatment of LNCaP human prostatic adenocarcinoma cells has been shown to induce several positive biomarkers for neuroendocrine differentiation including CgA<sup>(19)</sup>. We detected very few CgA- and NSE-positive cells in tumour tissue. The reduction of CgA-positive cells in tumour tissue corroborates earlier results from serum, showing a reduction in localised PCa and an increase in castration-resistant PCa<sup>(31)</sup>.

The ability of genistein to modulate the progression of existing PCa is not clear. In the present study, we have investigated the effects of pure genistein. As opposed to the majority of chemopreventive reports of genistein treatment, some recent studies on mouse xenograft models indicate genistein to promote increased metastasis of PCa, whereas isoflavones containing genistein and daidzein do not (33-37). Other studies show that dietary genistein inhibits metastases of human cancer, including PCa, in mice and that the discrepancy may be related to methodology<sup>(38,39)</sup>. In addition, Setchell et al.<sup>(40)</sup> raised doubts about the use of rodent models for gaining insight into the effect of isoflavones in humans due to differences in the metabolism of genistein. We did not detect any signs of cancer-promoting effects in our study with pure





B. Lazarevic et al.

genistein intervention in our human study subjects having early localised PCa.

Overall, genistein intervention at nutritionally relevant levels in patients with early PCa modulated several biomarkers which may be related to cancer prediction and progression. Genistein may have an inhibitory effect on androgen-related biomarkers. We also suggest a possible mechanism as to how genistein may induce cell cycle arrest and inhibition of proliferation in PCa. A limitation in our study is the small number of cases included and also the relative short time of intervention. Further studies on the effects of genistein in PCa are warranted, including clinical studies examining biomarkers and in vitro studies investigating its mechanisms of action.

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