

# SmartGenomics™ Prostate Profile

## Oncology Services

Clinical to Genomic

### → Advanced Standard of Care

**PathGroup SmartGenomics: Prostate** is designed for *use at diagnosis* of primary or metastatic prostate carcinoma to uncover therapeutic options and aid in treatment planning to improve patient outcomes.

- Clinically actionable genomic information for 44 gene mutations, cytogenetic abnormalities (FISH), and whole genome copy number changes (CMA) from a single biopsy
- Fully integrated testing on every case for a complete patient picture

### → Tailored Genomic Prostate Profile

#### Next Generation Sequencing (NGS)

AKT1	Therapeutic implications via PI3K/AKT/mTOR inhibition
APC	Therapeutic target, independent predictor of a poor prognosis in prostate cancer
ARAF	Potential therapeutic targets
ATM	Confers sensitivity to platinums and PARP inhibitors
BRAF	Potential therapeutic target
BRCA1/2	Confers sensitivity to platinums and PARP inhibitors
CDKN2A	Plays a critical role in prostate cancer progression, prognostic marker
CTNNB1	Potential therapeutic target
EGFR	Prognostic value in prostate cancer, potential therapeutic target
ERBB2/4	Related to metastatic potential of prostate cancer
FBXW7	Tumor suppressor linked to prognosis
FGFR1/2/3/4	Potential driver mutation and therapeutic target
IDH1/2	Deregulates cellular metabolism, potential therapeutic target
JAK1/2/3	Potential therapeutic targets
KDR	Recently discovered gene of importance in prostate cancer
KIT	Promotes migration and invasion of prostate cancer cells
KRAS	Potential therapeutic target
MET	Therapeutic target
NF1/2	Tumor suppressor function, potential prognostic value
NOTCH1	Deregulation is a feature of tumorigenesis
NRAS	Potential therapeutic target
NTRK1/2/3	Potential gene fusion partner, therapeutic target
PDGFRA	Potential therapeutic target
PIK3CA	Potential response to TORC1 inhibitor rapamycin and its analogues: everolimus, temsirolimus, and ridaforolimus
PTEN	Potential response to mTOR inhibitor everolimus, PI3K / AKT inhibitors under study
SMAD4	Suspected gatekeeper gene
SMO	Implicated in the development of PCa and progression to more advanced and castration-resistant disease
SRC	Potential therapeutic target
STK1	Tumor suppressor gene
TP53	Patients with TP53 mutations may have better outcomes with bevacizumab, therapeutic target
TSC1/2	Suspected gatekeeper gene
VHL	Potential regulator of PDL1 expression

(profile components listing continued on back)

### Fluorescence In Situ Hybridization (FISH)

PTEN	Prognostic and predictive in combination with ERG
AR	Provides therapeutic guidance
ERG	Prognostic and predictive in combination with PTEN

### Cytogenomic Microarray (CMA)

Whole genome copy number changes in >22,000 genes, 500 of which are implicated in cancer