

# SmartGenomics™ Thyroid Profile

## Oncology Services

### → Advanced Standard of Care

Clinical to Genomic

**PathGroup SmartGenomics: Thyroid** is designed to identify the likelihood of malignancy and provide prognostic and therapeutic information for indeterminate, suspicious, or malignant thyroid fine needle aspiration (FNA) as well as patients previously diagnosed with malignant thyroid cancer

- Clinically actionable genomic information for 21 genes, targetable thyroid carcinoma driver fusions, and PAX8 FISH analysis
- Full results in 7 to 10 days

### → Tailored Genomic Thyroid Profile

#### Next Generation Sequencing (NGS)

|         |  |
|---------|--|
| AKT1    | Advanced and dedifferentiating thyroid tumors; PI3K/AKT inhibitors under study   |
| ALK     | Anaplastic thyroid carcinoma, potential benefit from treatment with ALK inhibitors                                       |
| BRAF    | Commonly occurring mutation in thyroid carcinoma; potential benefit to RAF inhibitors                                    |
| CTNNB1  | Well differentiated thyroid cancer, poorly differentiated and anaplastic carcinomas; potential benefit to RAF inhibitors |
| E1F1AX  | Eukaryotic translation factor thought to be involved in tumorigenesis  |
| ERBB2/4 | Reported in undifferentiated thyroid cancer; potential benefit to TKIs   |
| GNAS    | Autonomously functioning benign thyroid nodules  |
| HRAS    | Papillary thyroid cancer; medullary thyroid carcinomas   |
| KRAS    | Papillary thyroid cancer; medullary thyroid carcinomas   |
| MET     | Papillary thyroid cancer, inherited Medullary Thyroid Cancer (MTC) and sporadic MTC                                      |
| NRAS    | Papillary thyroid cancer and medullary thyroid carcinomas; potential benefit to MEK inhibitors and other TKIs            |
| PIK3CA  | Advanced and dedifferentiating tumors and anaplastic thyroid cancers; PI3K/AKT under study                               |
| PTEN    | Seen in advanced and dedifferentiating tumors  |
| RET     | Sporadic and familial forms of medullary thyroid carcinoma; RET inhibitors in development                                |
| SMAD4   | Identified in papillary thyroid carcinoma proliferation  |
| SMO     | Follicular thyroid adenomas, anaplastic thyroid carcinomas, and papillary thyroid carcinomas                             |
| SRC     | Target of dasatinib in PTC   |
| TERT    | Associated with aggressive thyroid tumor characteristics, tumor recurrence and patient mortality                         |
| TP53    | Aggressive, well differentiated thyroid cancer; poorly differentiated and anaplastic carcinomas                          |
| TSHR    | Signaling gene known to be mutated in various thyroid carcinomas   |

#### Gene Fusions

Targetable driver fusions in various thyroid carcinomas:  
ALK, BRAF, FGFR1/2/3, MAML2, MET, NTRK1/2/3, PIK3CA, PPARG, RET, TERT

#### Fluorescence In Situ Hybridization (FISH)

PAX8 A reliable means of discerning thyroid origin for undifferentiated tumors of the head and neck

# SmartGenomics

## THERANOSTIC SUMMARY REPORT

Original Procedure Date: 05/15/2013, Sample Type: Right Thyroid, Block Information: A6, Original Accession number:  
Tumor Percentage: 81-100%  
Original Diagnosis: Carcinoma-Thyroid Origin, Original Diagnosis rendered at:

Clear, concise results

### RESULTS

An **NRAS Q61K** was detected by targeted next generation sequencing.  
An **ABNORMAL** cytogenomic array result was detected.

### INTERPRETATIVE SUMMARY

Review of the accompanying pathology report indicates a history of follicular thyroid carcinoma. Microscopic examination confirms poorly differentiated carcinoma corresponding to the above referenced accession number. Sufficient tumor tissue is present for molecular analysis. Excellent gene coverage was achieved which passed our internal quality checks. **An NRAS Q61K mutation was detected by targeted next generation sequencing** interrogating the 35 genes summarized below. No other mutations were found in hot-spot regions of the other genes analyzed. Three different human RAS genes have been identified: KRAS, NRAS, and HRAS. The different RAS genes are highly homologous but functionally distinct. RAS proteins are small GTPases which cycle between inactive guanosine diphosphate (GDP)-bound and active guanosine triphosphate (GTP)-bound forms. RAS proteins are central mediators downstream of growth factor receptor signaling and therefore are critical for cell proliferation, survival, and differentiation. RAS can activate several downstream effectors, including the PI3K-AKT-mTOR pathway, which is involved in cell survival, and the RAS-RAF-MEK-ERK pathway, which is involved in cell proliferation. RAS has been implicated in the pathogenesis of several cancers. Activating mutations within the RAS gene result in constitutive activation of the RAS GTPase, even in the absence of growth factor signaling. The result is a sustained proliferation signal within the cell. Specific RAS genes are recurrently mutated in different malignancies. The frequency of NRAS mutations in thyroid carcinomas is 6%. While most non-thyroid cancers have mutations in KRAS codons 12 and 13, most thyroid tumors have been found to have mutations in NRAS codon 61 and HRAS codon 61. In the majority of cases, these mutations are missense mutations that introduce an amino acid substitution at position 61, as was observed in this case. The result of these mutations is constitutive activation of NRAS signaling pathways. Currently, there are no direct anti-NRAS therapies available, but preclinical models suggest that MEK inhibitors such as selumetinib and trametinib may be effective. A number of therapies are available in an investigational setting that target this genomic aberration. An example of clinical trial can be found below. Please also see below for a summary of single nucleotide polymorphisms (SNPs).

Therapeutic implications

### THERAPEUTIC ASSOCIATIONS

| Potential Therapeutic Response / Drug Class   | Disease Association | Gene / Locus | Alteration |
|---|---------------------|--------------|------------|
|  Investigational agents (i.e. MEK and PI3K inhibitors) may be available. | Various cancers     | NRAS         | Mutations  |

Personalized clinical trials

### CLINICAL TRIALS

| NCTID       | Title   | Conditions   | Location   | Sponsor                         |
|-------------|---|--|--|---------------------------------|
| NCT01529593 | Temsirolimus in Combination With Metformin in Patients With Advanced Cancers                          | Advanced Cancers   | UT MD Anderson Cancer Center<br>Houston, Texas, 77030, United States   | M.D. Anderson Cancer Center     |
| NCT01208051 | Cediranib Maleate With or Without Lenalidomide in Treating Patients With Thyroid Cancer               | Recurrent Thyroid Cancer, Stage I<br>Follicular Thyroid Cancer, Stage I<br>Papillary Thyroid Cancer, Stage II<br>Follicular Thyroid Cancer, Stage II<br>Papillary Thyroid Cancer, Stage III<br>Follicular Thyroid Cancer, Stage III<br>Papillary Thyroid Cancer, Stage IVA<br>Follicular Thyroid Cancer, Stage IVA<br>Papillary Thyroid Cancer, Stage IVB<br>Follicular Thyroid Cancer, Stage IVB<br>Papillary Thyroid Cancer, Stage IVC<br>Follicular Thyroid Cancer, Stage IVC<br>Papillary Thyroid Cancer | Multiple locations in United States, Canada  | National Cancer Institute (NCI) |
| NCT01182285 | A Phase II Trial of Valproic Acid in Patients With Advanced Thyroid Cancers of Follicular Cell Origin | Thyroid Neoplasm   | National Institutes of Health Clinical Center, 9000 Rockville Pike<br>Bethesda, Maryland, 20892, United States | National Cancer Institute (NCI) |