


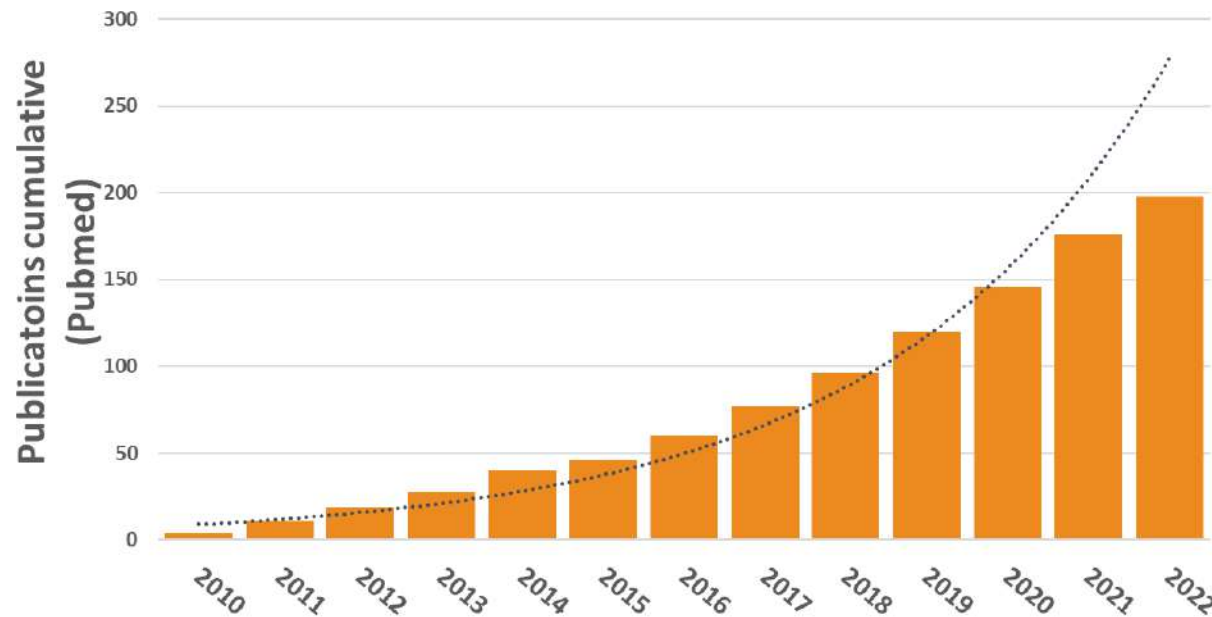
Introducing an unbiased approach that uses Multiplex Kinome Activity Profiling

Tracking changes in cellular signaling and pathways with PamGene



Pamgene overview

- Spin-off from the Organon Teknika 
- Contract Research Services and platform technology
- Specialized in signaling protein biomarker technology
- Extensive body of peer reviewed publications on technology
- Patented multiplex peptide biomarker profiling technology (15 patents)



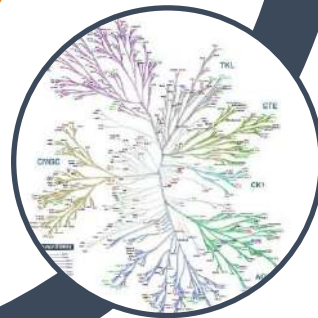
Trial overview



Direct access to our approach during this trial period of typically 12 months

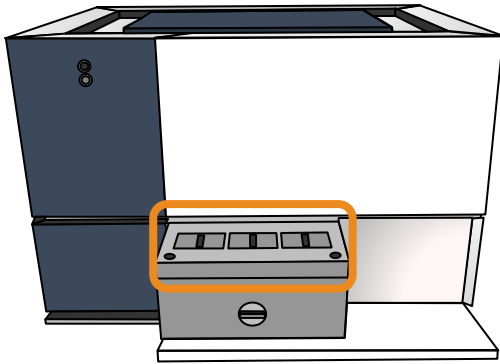


Multiple groups invest together and collaborate using the approach and incorporate us in new funded projects



**The investment of consumables is including our direct scientific support.
E.g.: Experimental design, data analysis, reporting, interpretation**

PamStation 12[®]



Reader instrument

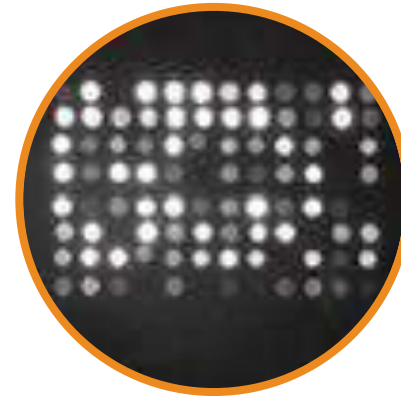
- Capable of loading 3 Pamchips per experiment, containing 12 samples
- High throughput, only 15 min's hands on time

Pamchip 4[®]



Consumables

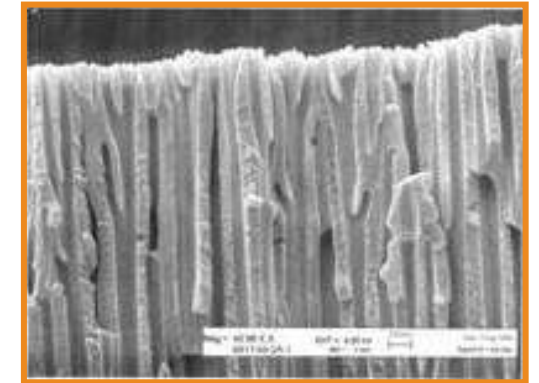
- Serine /Threonine Pamchip (STK) with 144 targets / substrates
- Tyrosine Pamchip (PTK) with 196 targets / substrates



One of 4 Arrays

- All arrays are identical
- Generic antibody binding to all phosphorylated targets / substrates

3D membrane



Porous structure

- Peptides are spotted in high concentrations, making the assay sensitive.
- During incubation, the lysate is pumped back and forth through the membrane, this makes the assay fast.

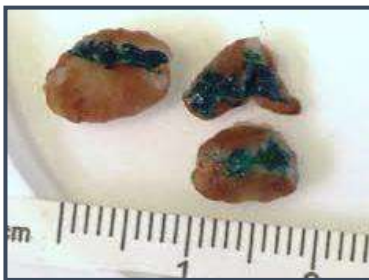
Input requirements

0,5 – 5,0 μ g total protein per array
10.000 – 50.000 cells per array



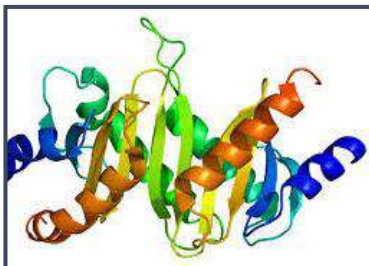
Primary and cultured cells

PBMCs, WBC, platelets, Bone marrow
Primary cells
Culture cells (adherent or suspension)



Primary biopsies, slices, clinical samples

Freshly frozen (alternatively Tissue-Tek)
Tumor content advise >70%
Different tissues (FNA) e.g. Colon, lung, liver, breast,
brain, prostate, skin, thyroid, CSF

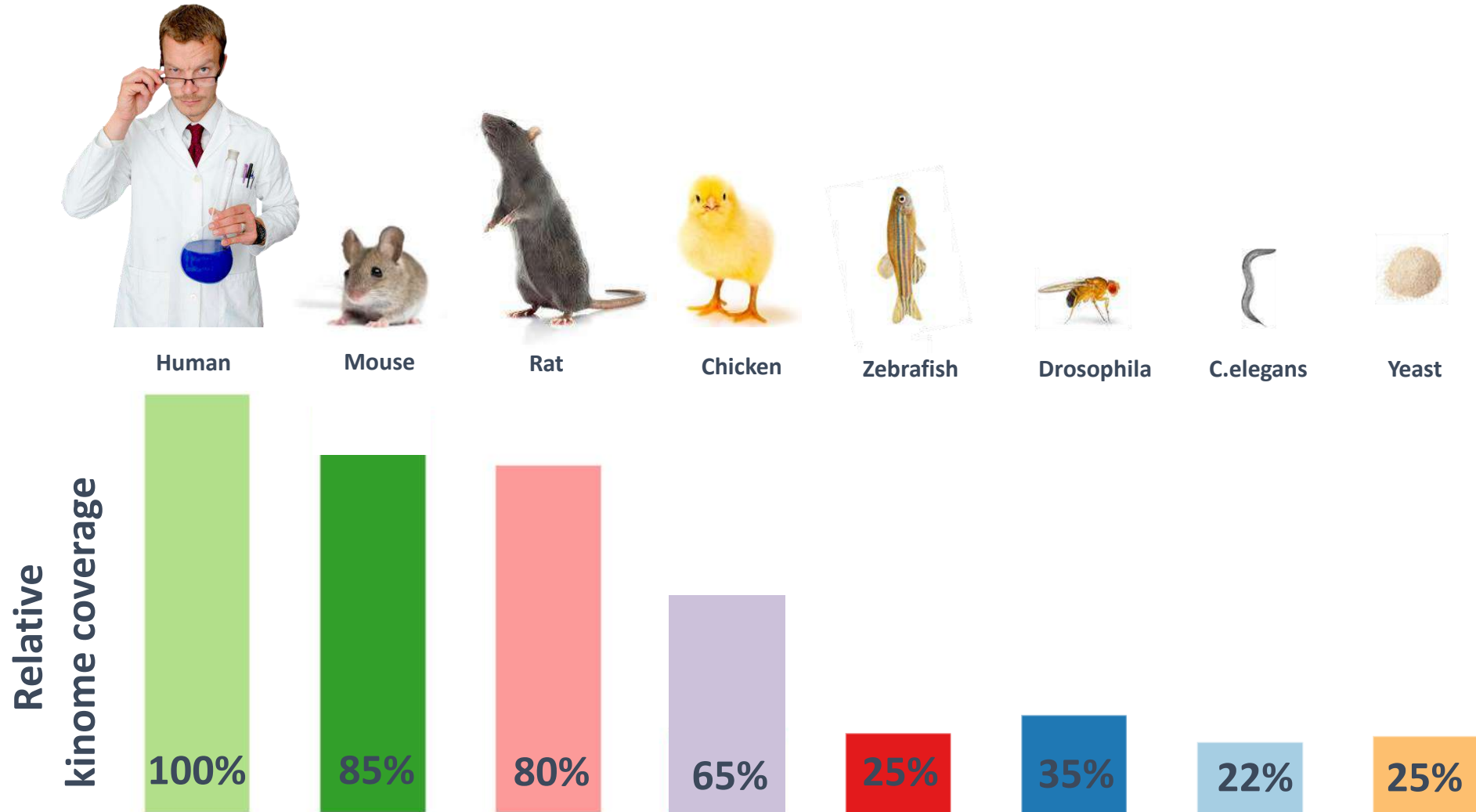


Purified proteins

Recombinant kinases

**Collect samples
over time, fresh
freeze, and store
for long periods
at -80°C**

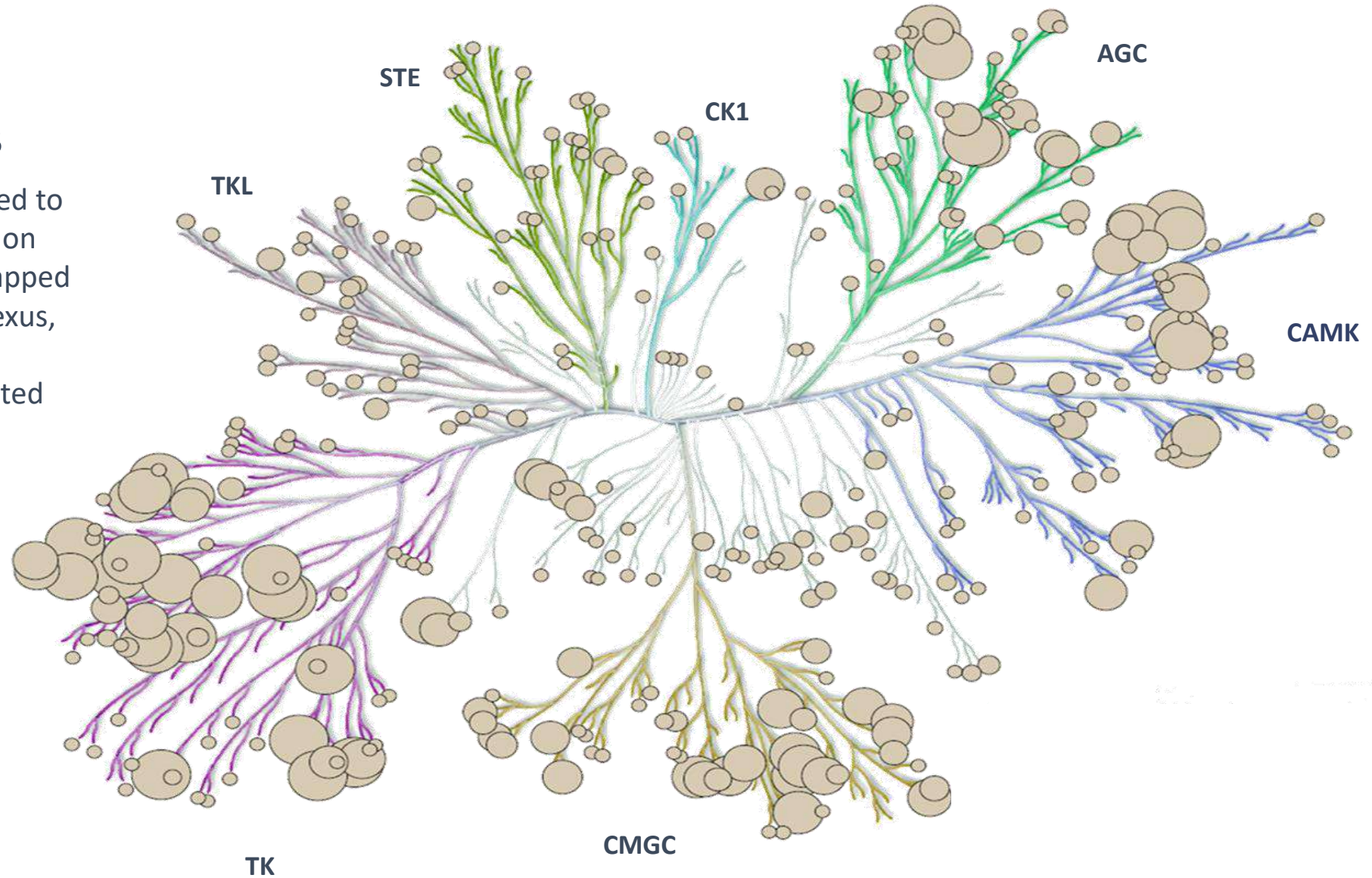
Organism coverage



Kinome Coverage

Covering >380 Kinases

Kinases that are known or predicted to phosphorylate substrate peptides on the PTK and STK PamChip® are mapped from select databases (HPRD, Kinexus, Phosphosite PLUS, Reactome, Phospho.elm, UniPROT) and projected on the kinome tree.



Applications



Fundamental and Discovery research

- Signaling & Pathway elucidation
- Target discovery
- Target interaction
- Disease model characterization

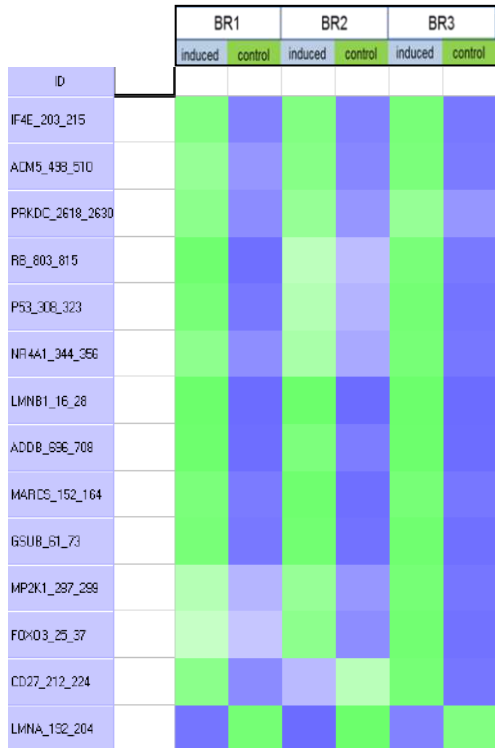
Biomarker and Clinical research

- Therapy-predictive biomarkers
- Prognostic biomarkers
- Classification biomarkers

Pamchip reveals increased MAPK signaling

Overexpression of Raf-1 is expected to result in an increase of ERK (MAPK) signaling

Significantly different phosphosites between induce en control

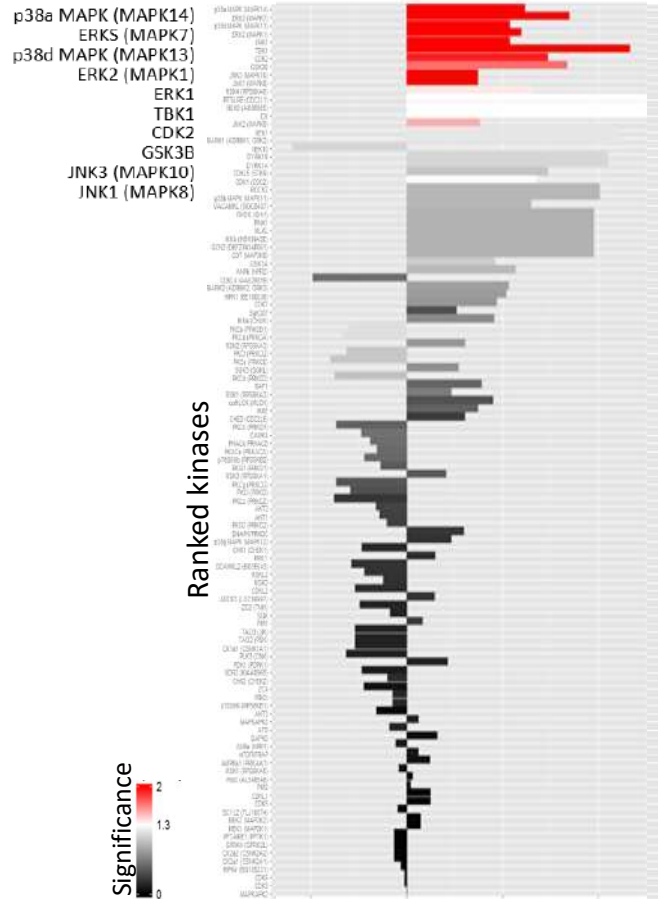


Higher phosphorylation

Lower phosphorylation



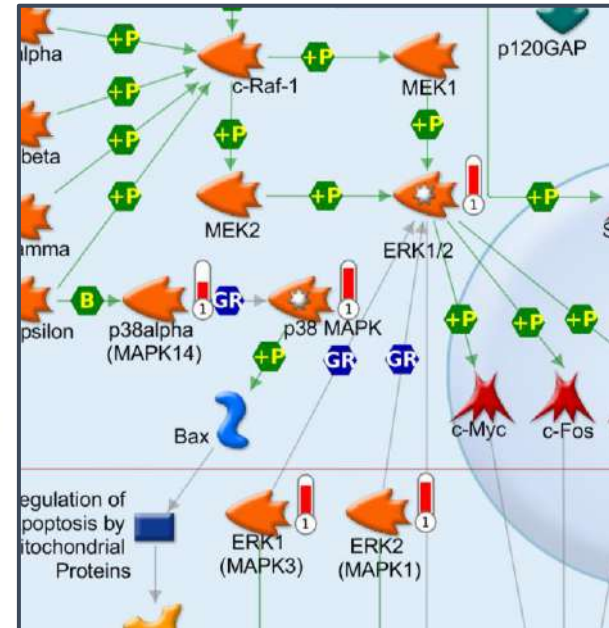
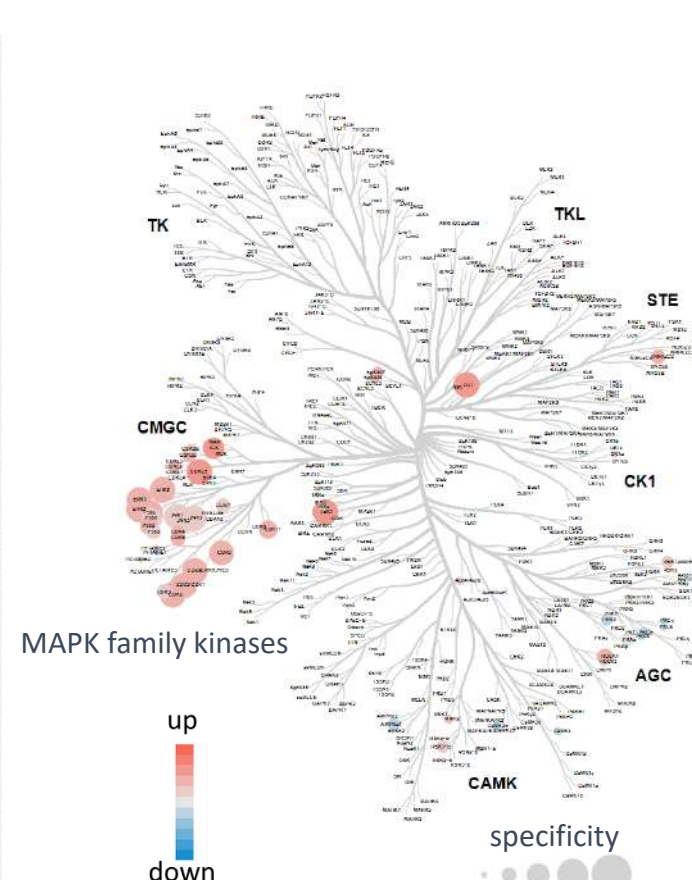
Upstream kinase analysis, based on Kinase to substrate interaction recorded in know databases



Significance



Upstream kinase analysis visualized on a kinome phylogenetic tree and Metacore pathway analysis

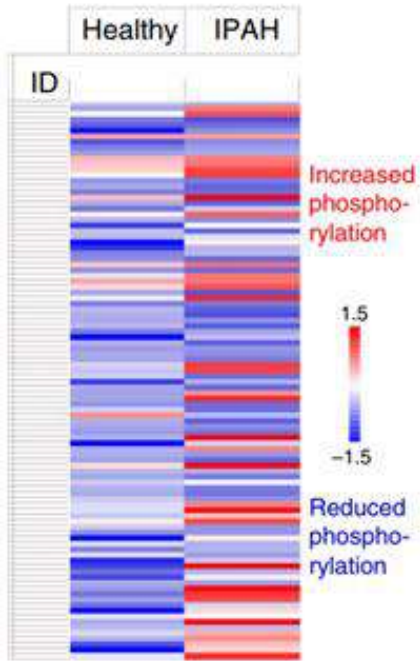


Pamchip reveals increased activity of the CDK signaling pathway

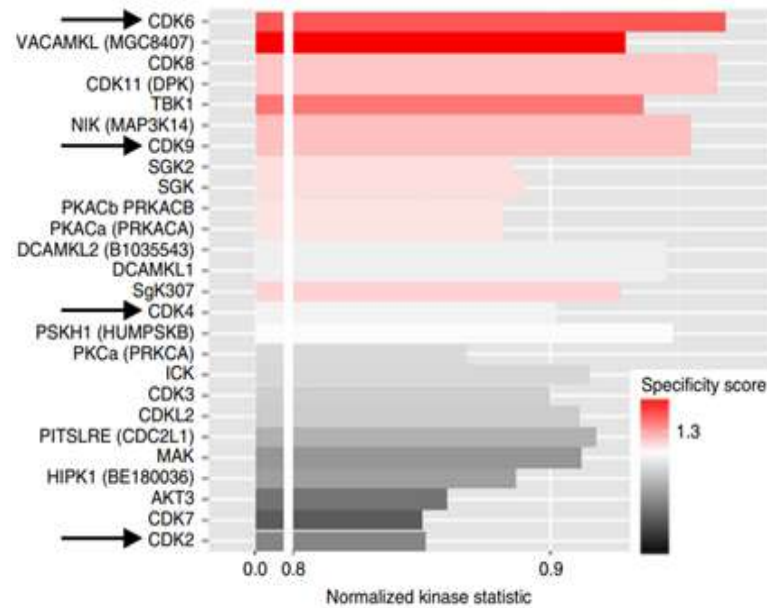
Profiling of primary pulmonary cells from healthy and IPAH patients

Validation

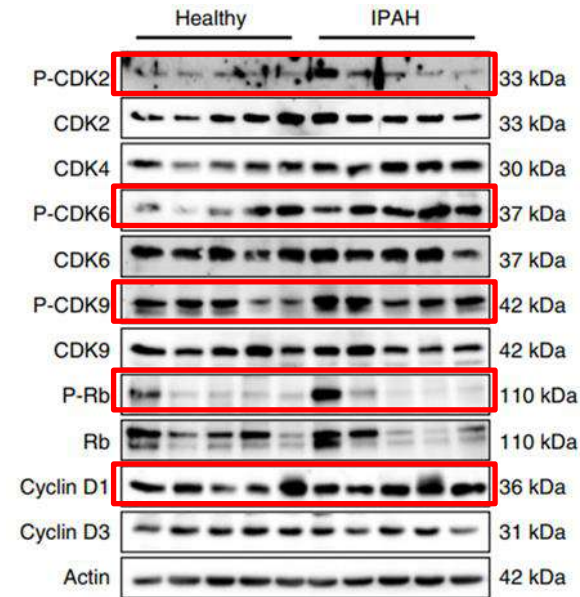
Differential peptide analysis



Upstream Kinase Analysis

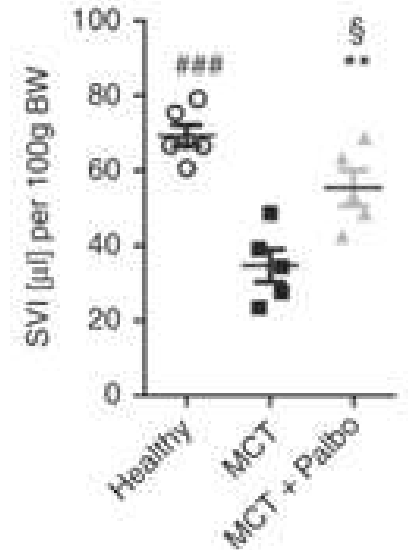


Western blot



In vivo proof MCT rat Model

Stroke volume index



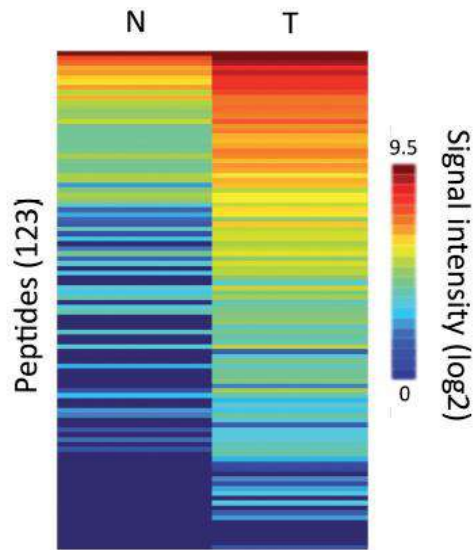
IPAH: Idiopathic pulmonary arterial hypertension

MCT = Disease inducer
Palbo.= CDK inhibitor

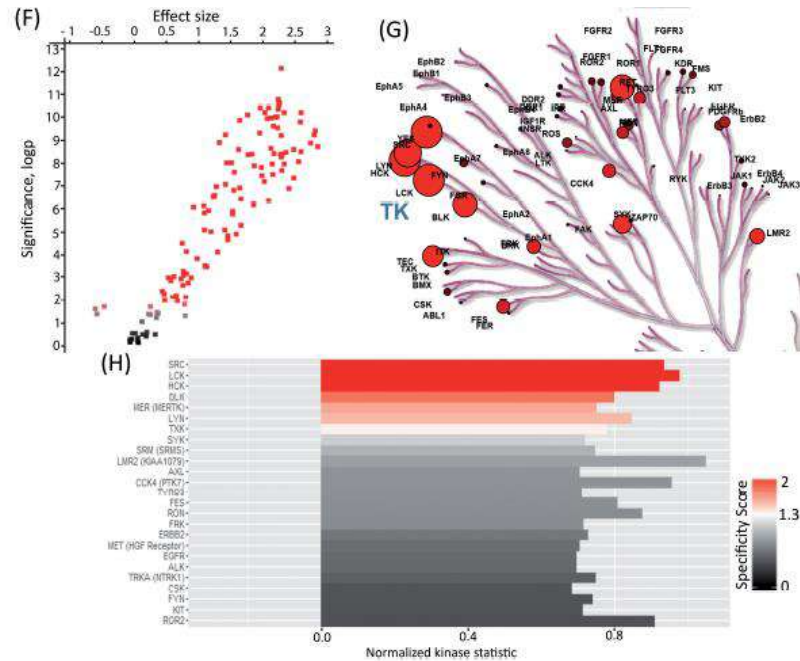
Increased activity of Src family signaling in HNSCC tumours

Profiling of tumour vs healthy tissues in HNSCC patients

Differential peptide analysis



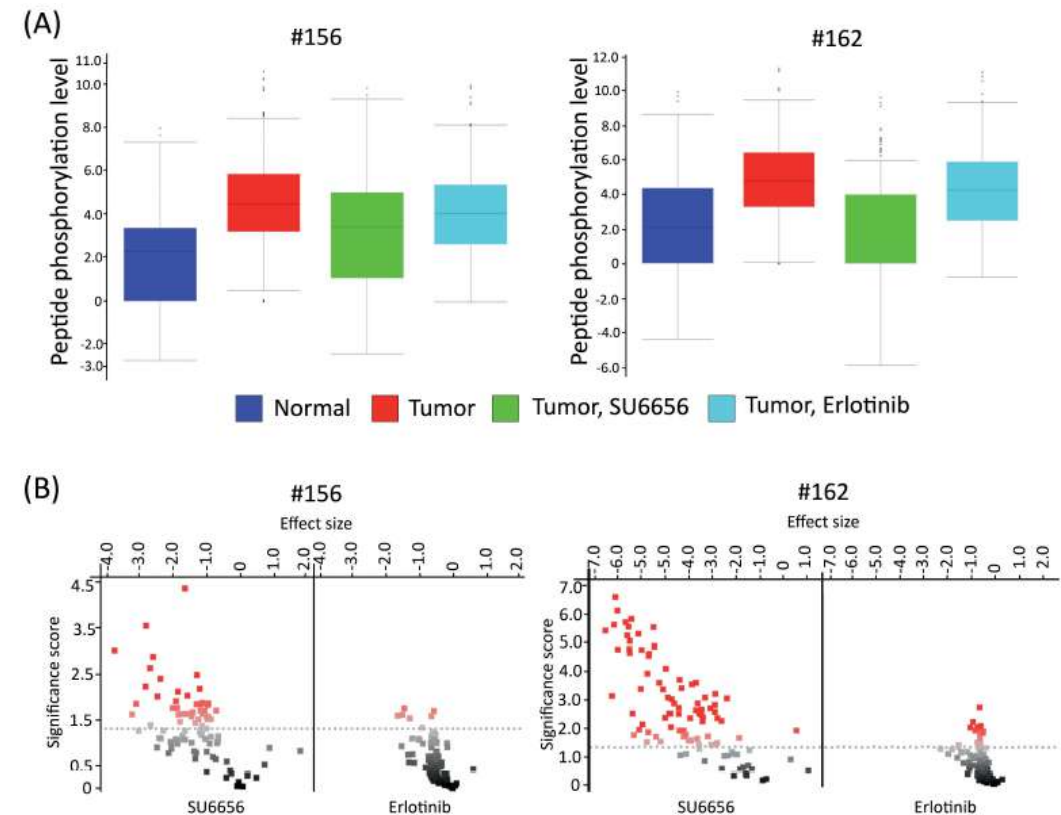
PamGene PTK Analysis



HNSCC, head and neck squamous cell carcinoma

Validation

Ex vivo inhibition of patient lysates with SFK inhibitor SU6656



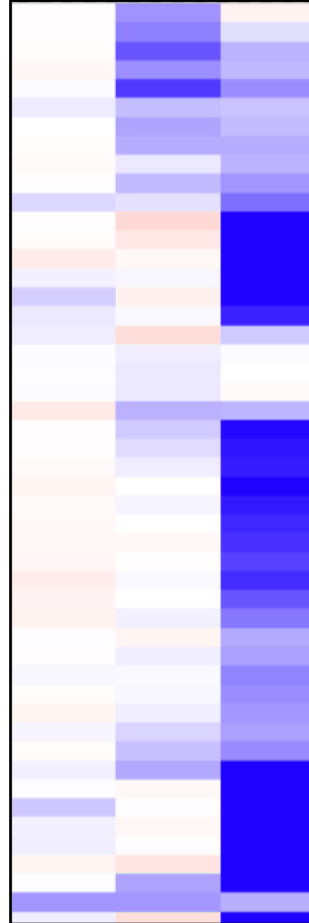
Directly adding kinase inhibitors in lysate

Cellular Treatment

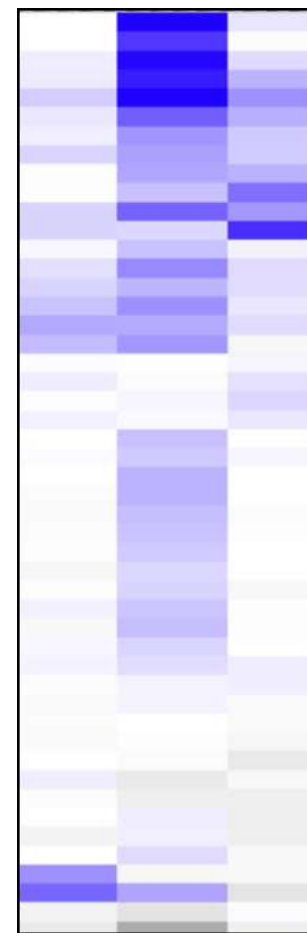


Also non-kinase inhibitors (HDAC inhibitors) can be profiled on their indirect effect on kinases in cells.

DMSO MTKI HDAC



DMSO MTKI HDAC



Lysate Treatment

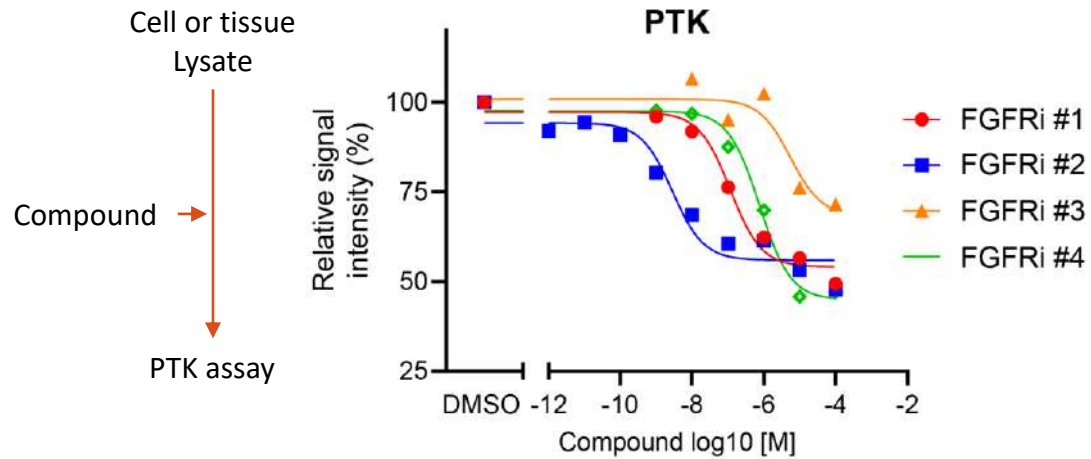


4-fold inhib.



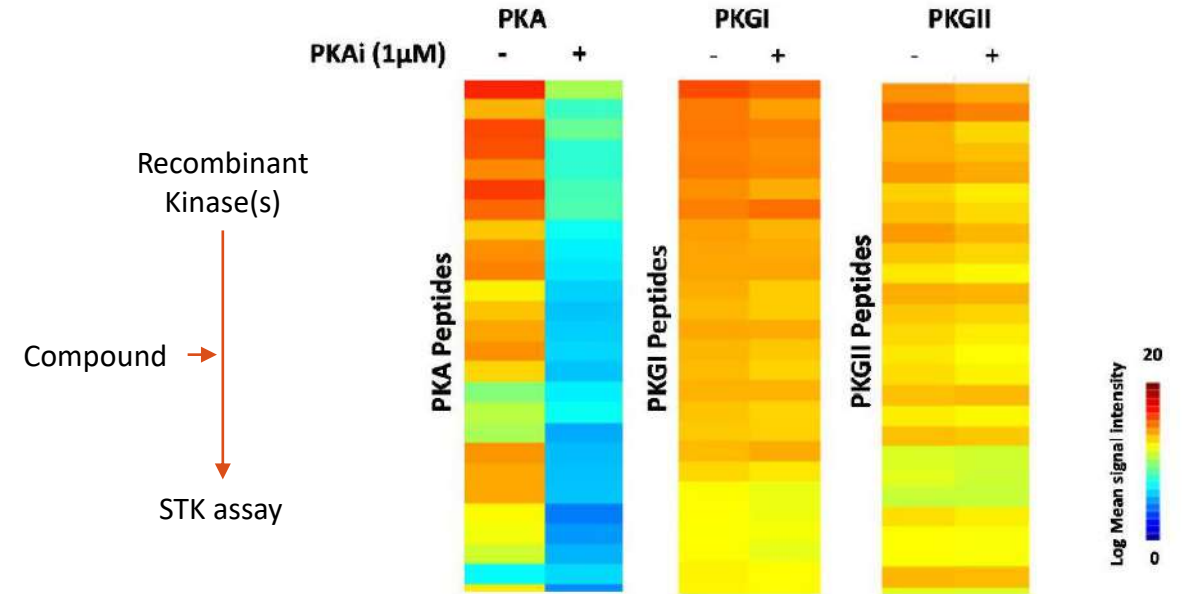
No inhib.

In vitro IC₅₀ determination of novel FGFR inhibitors in specific cell lysates



Courtesy of Services

Specificity of PKA inhibitor on recombinant PKA and PKG



Roy et al., *Int. J. Mol. Sci.*, 2021

Consumable investment



A single Trial Kit Purchase of € 4.000,00 euro.

- **Each kit contains a single type of 12 PamChips, separately sealed. That translates to In total 48 identical arrays (STK or PTK), to investigate 48 samples Necessary reagents are included**
- **This investment includes:
The installation of the instrument on location;
Training of users in instrument handling & wet lab;
We perform data analysis, shared in personalised reports;
Scientific support for preparation and interpretation.**
- **To allow us to start, we need enough interest on a specific location, typically that translates to 5 - 8 groups that invest 10-16 kits for their research purposes.**



Reach out at: jlebens@pamgene.com

Pamgene IVD certification June 2022

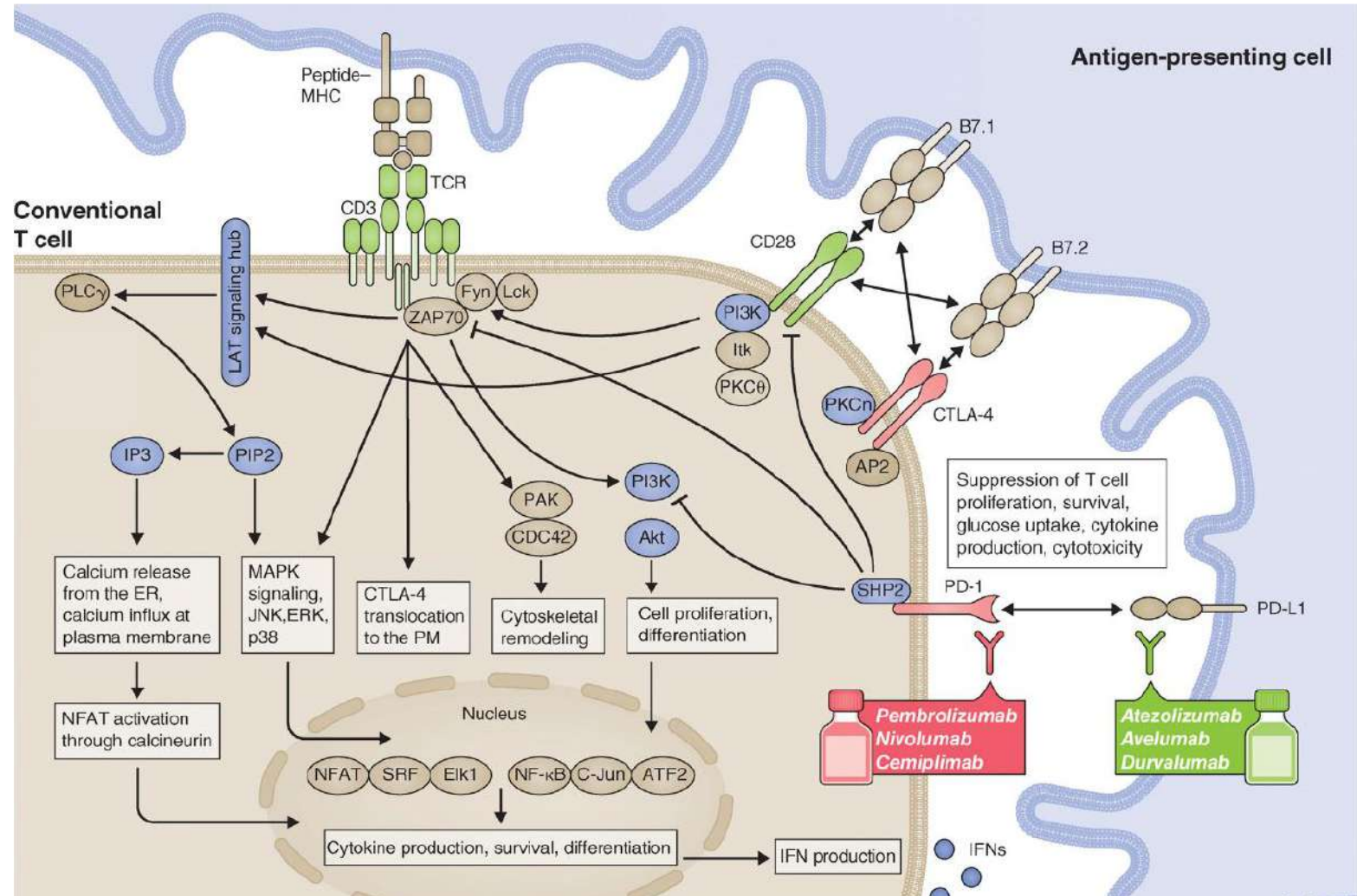
A quick Introduction to our internal IVD development program



Diagnostic Application

Kinase activity impacts how individual patients respond to PD-1 and CTLA-4-directed therapies.

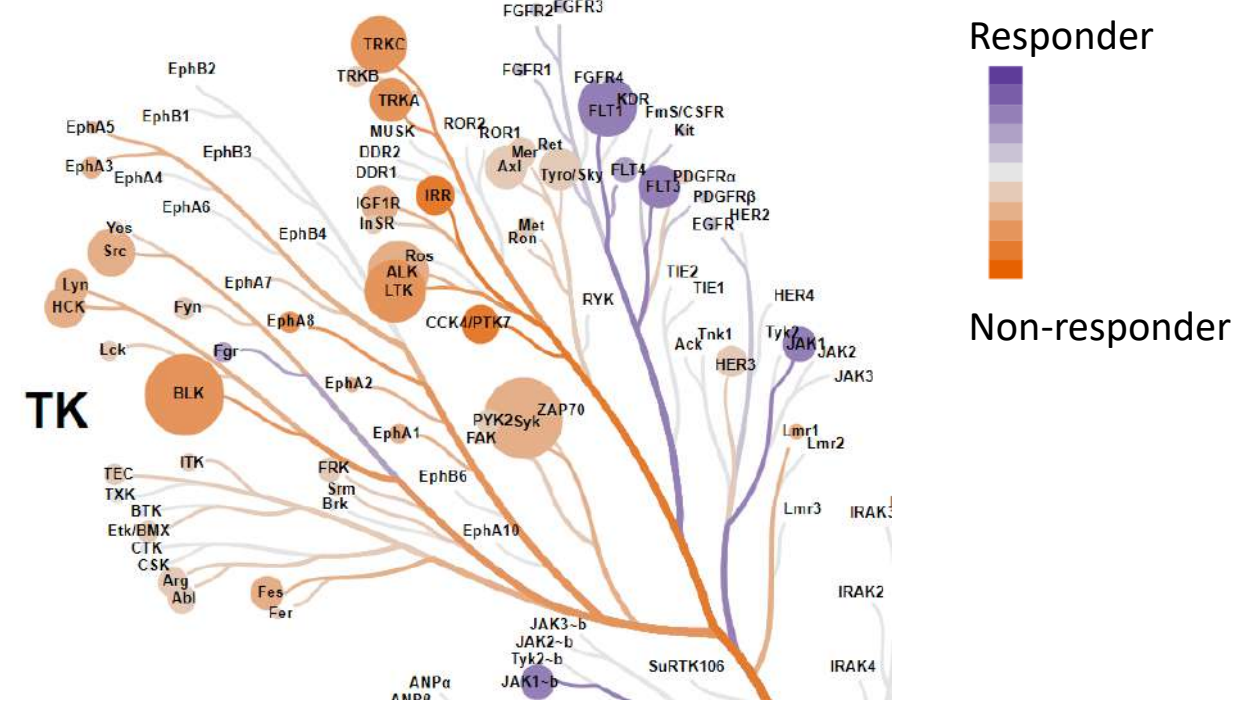
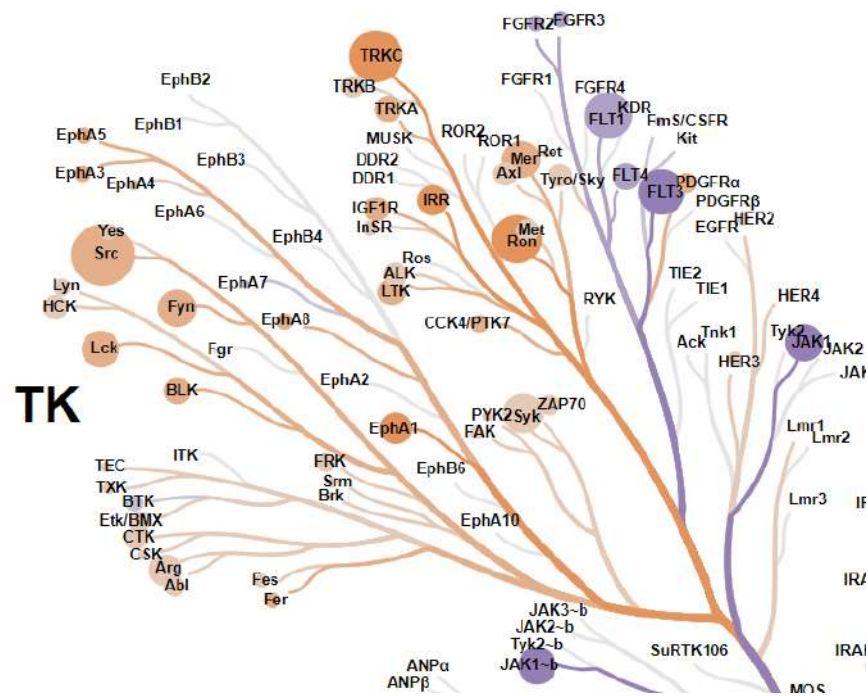
Our diagnostic response prediction is done by profiling whole blood PBMC samples, taken from NSCLC & Melanoma patients prior to treatment.



Differential kinases between responders and non-responders

Melanoma Cohort
PD20 / Clinical calibration

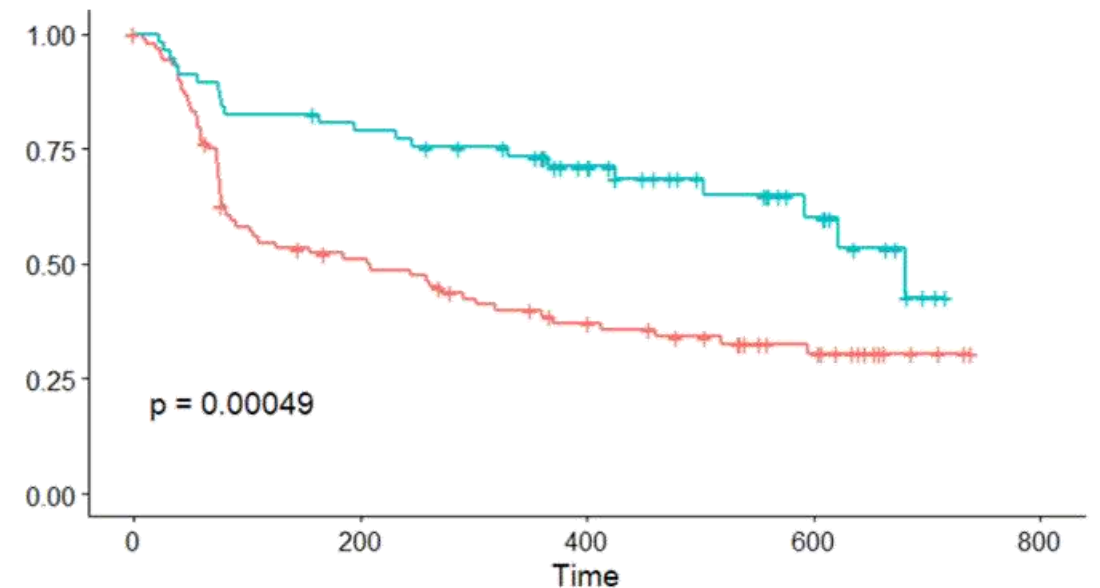
NSCLC Cohort
PD12 / clinical calibration



Prediction performance in Melanoma validation study

- Study design: 160 patients with advanced melanoma and eligible for ICI treatment (Standard of care), multi-center study.
- Prospective trial, 5 clinical centers (calibration & validation)
- Two cohorts (160 samples)have been analysed:
 - Centralized, standardized PMBC isolation.
 - Samples analysed in Diagnostic Assay Services Facility.
 - Kinome multiplex array data analysed
- Patient response is assessed by DR (durable response) and PFS (Progression Free Survival).
- Accuracy in independent validation cohort is **70-76 %** for immune treatments
- Hazard Ration(HR) for the survival analysis ranges from 1.4 to 2.1
- Multimodal analysis ongoing (clinical chemistry, hematology)
- **IOpener**® test is predictive for disease progression in Melanoma patients treated with anti-PD1. (first line and/or second line immune treatment)

Melanoma validation studies
Combined cohorts -> DLM model

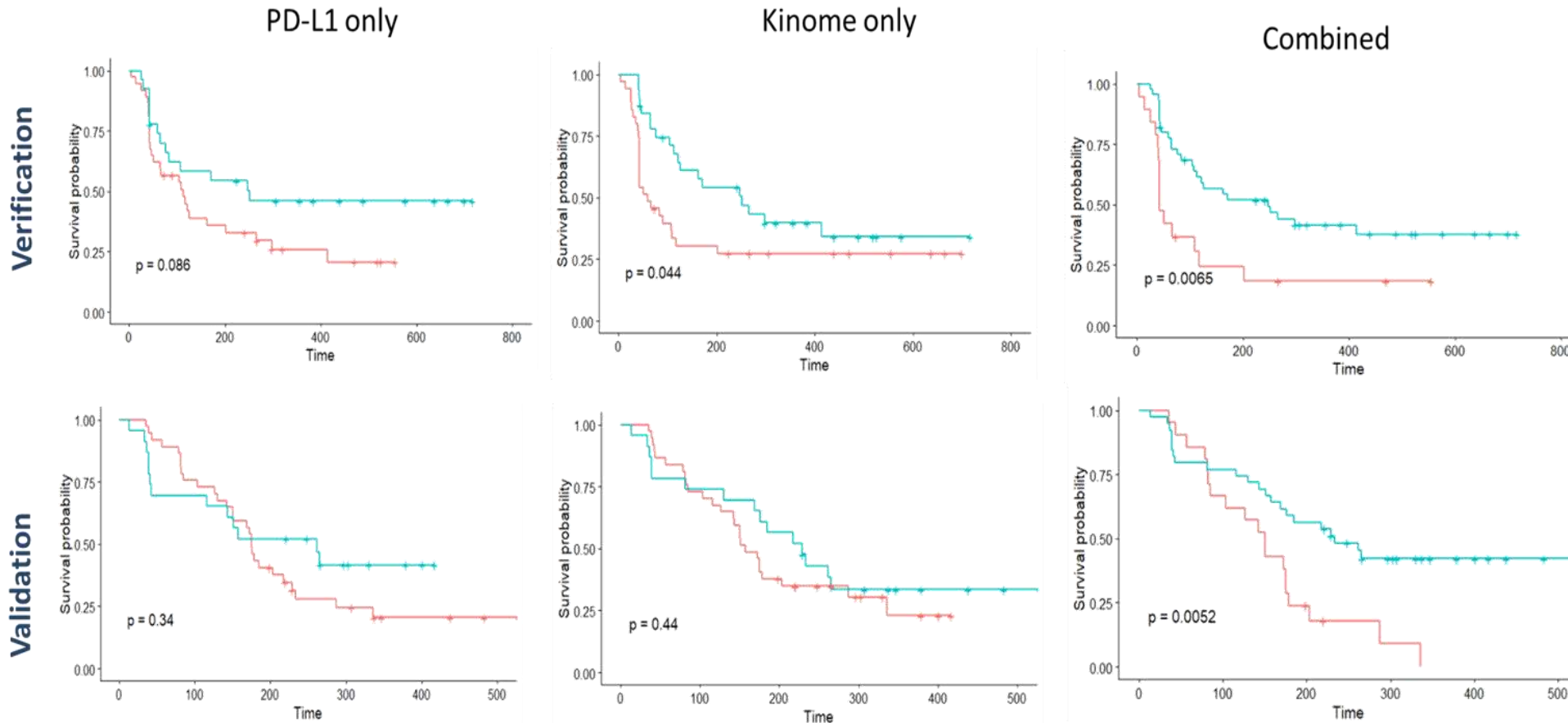


Predicted responder

Predicted no responder

Prediction performance in NSCLC validation study

- True and false positive/negative responders are assessed for PD-L1 and **IOpener**[®] prediction score in NSCLC validation studies (n=150)
- **IOpener**[®] in combination with PD-L1 test is superior predictor of the response of NSCLC patients treated with ICI therapy
- Hazard Ratio (HR) for TPS score is 1.3 and HR is 2.3 when combined with IOpener (Accuracy is 72% for TPS<50%)
- **IOpener**[®] test as an inclusion criterion will significantly improve the response to combo-therapy (TPS < 50%) compared to mono therapy (TPS>50%).



Predicted responder

Predicted no responder

A view on our customers



Pharmaceutical Companies
Biotechnology Companies
Academic Medical Centers
Research Institutes
Laboratory Services Providers

