

# Analysis of active oncogenic signal transduction pathways in HGS ovarian cancer

## HH pathway activity associated with platin resistance

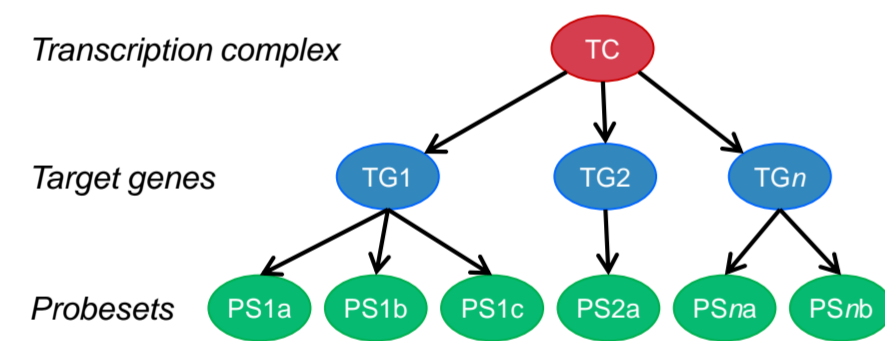
Paul van de Wiel<sup>1</sup>, Jozien Helleman<sup>2</sup>, Eveline den Biezen<sup>1</sup>, Janneke Wrobel<sup>1</sup>, Henk van Ooijen<sup>1</sup>, Wim Verhaegh<sup>1</sup>, Lena van Doorn<sup>2</sup>, Els Berns<sup>2</sup>, Anja van de Stolpe<sup>1</sup>,  
(1) Precision Diagnostics, Philips Research, Eindhoven, The Netherlands; (2) Dpt of Medical Oncology, ErasmusMC, Rotterdam, The Netherlands

### Introduction

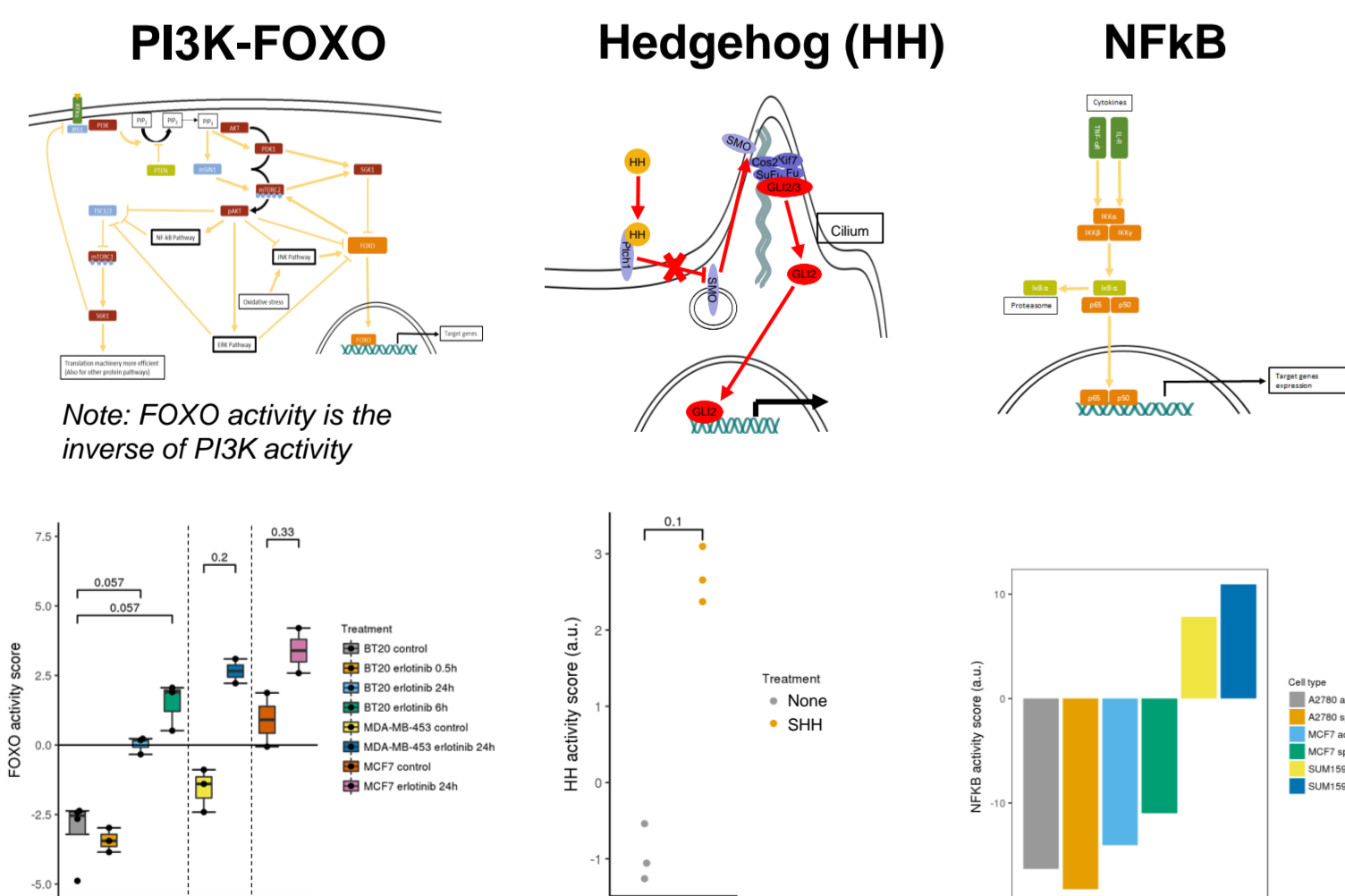
Patients with ovarian cancer in general have a bad prognosis due to frequently advanced disease at diagnosis associated with limited therapeutic options, and recurring disease. Use of targeted drugs in ovarian cancer is not very advanced, one major reason being insufficient knowledge on tumor driving signaling pathways. Hence there is a strong need for a better diagnostic approach.

### Signal transduction pathway analysis approach

A new diagnostic approach is being developed [Verhaegh et al. Cancer Res 2017;74(11):2936-45] that enables quantitative measurements of the functional activity status of (oncogenic) signal transduction pathways in cell and tissue samples, based on computational inference of pathway activity from measurements of mRNA levels of well-validated direct target genes of the transcription factor associated with the respective signalling pathway. mRNA levels can be measured using Affymetrix HG-U133Plus2.0 or +PM microarrays or RT-qPCR.



### Validation examples of pathway models



GSE30516: Erlotinib (EGFR inhibitor) inhibits PI3K pathway activity  
 GSE29316: Sonic Hedgehog induces HH pathway activity in fibroblasts  
 GSE43657: NFkB activity in inflammatory breast cancer cell line SUM159

### Materials and methods

Affymetrix HG-U133+PM microarrays were run on fresh frozen samples from 37 primary high grade serous (HGS) ovarian cancer patients (90 samples in total, from multiple metastatic locations). These data, and data from a number of publicly available Affymetrix datasets were analysed for signal transduction pathway activity, including Hedgehog (HH), PI3K and NFkB pathway activity.

### Results

#### HGS ovarian cancer patient samples, Erasmus MC

Two common pathway patterns were found:

- NFkB+FOXO active, i.e. PI3K inactive
- PI3K+Hedgehog (HH) active

Patient 17, primary carboplatin/paclitaxel resistant HGS ovarian cancer, with rapid disease progression and death: HH active and PI3K active (FOXO inactive)

Pt no	stage	FIGO	FIGO (prev. debulk, surg.)	Chemotherapy	IDS (interval debulking surgery after)	PD event	Time to PD (months)	Time to death (months)	Loc	%	O	E	NE	PI3K (inact)	NFkB act	HH act
1	3	IC	yes, incomplete	3x CisPT + 2x CT	no	1	36.6	1	53.9	Adnex left	80	12.9	3.9	2.5	X	
2	3	IIIc	yes, incomplete	CT	no	1	36.6	1	53.9	Ovarium	90	13.2	5.1	10.6		
3	3	IIIc	yes, incomplete	CisPT/ 5x 2x CT	yes, rest < 1 cm	1	36.3	1	45.6	Omentum	30	15.4	1.6	3.7	X	
4	3	IIIc	yes, incomplete	3x CisPT	no	0	302.0	1	302.0	Ovarium right	70	12.6	1.7	8.9	X	
5	3	IIIc	yes, incomplete	CT/ 6x 3w	no	1	34.0	1	81.7	Ovarium	45	13.3	3.2	12.3	X	
6	3	IV	yes, incomplete	CT/ 5x 3w	yes, complete	1	18.0	1	40.7	Ovarium	100	14.8	4.4	14.6	X	
7	2	IIb	yes, suboptimal	CT/ 3x	no	1	50.6	1	25.8	Omentum	40	7.3	5.8	11.9	X	
8	3	IIIc	yes, incomplete	6x CT	no	1	31.3	1	126.1	Ovarium	70	10.8	3.3	7.9	X	
9	3	IIIc	yes, incomplete	3x CT/ 3x paclitaxel/etoposide	yes, complete	1	21.7	1	21.7	Ovarium	90	3.9	3.8	8.3		
10	3	IIIc	yes, optimal	CT	no	1	11.5	1	13.5	Omentum	70	6.9	6.0	6.8	X	
11	IIIc	IIIc	yes, optimal	6x CT	no	0	133.8	0	133.8	Omentum	80	12.9	0.3	6.8	X	
12	3	IV	yes, optimal	6x CT	no	1	37.8	1	41.2	Ovarium left	70	10.8	4.4	12.0	X	
13	1-2	IIIc	yes, incomplete	6x CT + 3x CT	yes, optimal	1	21.0	1	84.7	Ovarium	80	6.2	1.9	12.4	X	
14	3	IIIc	yes, optimal	9x CT/ 3w	no	1	27.4	1	61.8	Adnex left	70	8.3	4.2	6.0	X	
15	2	IIIc	yes, optimal	6x CT	no	1	77.5	0	103.9	Ovarium	70	2.7	3.8	2.3	X	
16	2	IIIb	yes, optimal	no	no	1	1.2	1	1.2	Omentum	70	2.1	7.0	2.3		
17	3	IIIc	yes, complete	6x CT	no	1	6.5	1	11.1	Omentum	90	4.0	3.5	11.8	X	
18	2	IIIc	yes, suboptimal	9x CT	yes, complete	1	14.4	1	69.9	Ovarium	90	4.4	4.8	8.7	X	
19	IIIc	IV	yes, incomplete	3x 1x CT	yes, complete	1	69.2	1	75.1	Omentum	50	7.8	6.6	16.1	X	
20	2	IIIc	yes, incomplete	6x w CT + 4x CT then IDS	yes, complete	1	7.5	1	8.1	Ovarium	70	7.9	6.5	5.4	X	
21	3	IIIb	yes, complete	CT	no	1	20.2	1	20.2	Ovarium	60	11.6	1.8	10.4	X	
22	3	IIIc	yes, complete	CT	no	1	25.6	1	41.4	Ovarium	90	4.1	2.5	12.7	X	
23	3	IIIc	yes, complete	CT	no	1	25.6	1	41.4	Ovarium	90	4.1	2.5	12.7	X	
24	3	IIIb	yes, incomplete	3x CT + 6x CT	yes, complete	1	20.3	1	78.7	Omentum	80	5.8	5.6	13.3	X	
25	2	IV	yes, incomplete	6x w CT	no	1	2.0	1	2.0	Ovarium left	80	5.8	4.5	6.5	X	
26	3	IIIc	yes, optimal	9x CT	no	1	37.7	1	50.3	Ovarium	50	20.1	1.9	11.8	X	
27	3	IIIc	yes, complete	6x w CT	no	1	83.5	0	76.1	Adnex left	50	10.5	3.1	6.9	X	
28	3	IV	yes, incomplete	6x w CT + 5x CT + Tarceva (erlotinib)	no	1	16.0	1	12.2	Ovarium	60	11.9	1.1	12.7	X	
29	2	IIIc	yes, incomplete	6x w CT + 6x CT	no	1	19.0	1	65.6	Omentum	70	7.6	11.0	3.9	X	
30	2	IV	yes, optimal	9x CT	no	1	29.0	1	52.5	Omentum	60	10.3	5.1	9.2	X	
31	2	IIIc	yes, complete	no referral pat	no	1	131.6	0	232.2	Adnex	70	16.4	4.2	5.5	X	
32	3	IIIc	yes, incomplete	3x CT + 3x CT after IDS	yes	0	73.2	0	73.2	Ovarium right	90	12.4	0.4	9.8	X	
33	3	IIIc	yes, optimal	CT	no	1	11.2	1	37.4	Ovarium	80	11.8	4.7	16.3	X	
34	IIIc	IIIc	yes, complete	CT	no	1	11.2	1	37.4	Ovarium	80	11.8	4.7	16.3	X	
35	2	IIIc	yes, suboptimal	7x w CT + 2x CT	yes, complete	1	28.5	1	28.8	Omentum	50	17.7	1.6	13.3	X	
36	3	IIIc	yes, suboptimal	6x CT	no	1	54.2	0	71.8	Omentum	45	10.5	4.1	20.9	X	
37	3	IIIc	yes, incomplete	3x CT + 2x CT after IDS	yes, complete	1	20.7	1	63.3	Ovarium	80	11.7	6.7	6.8	X	

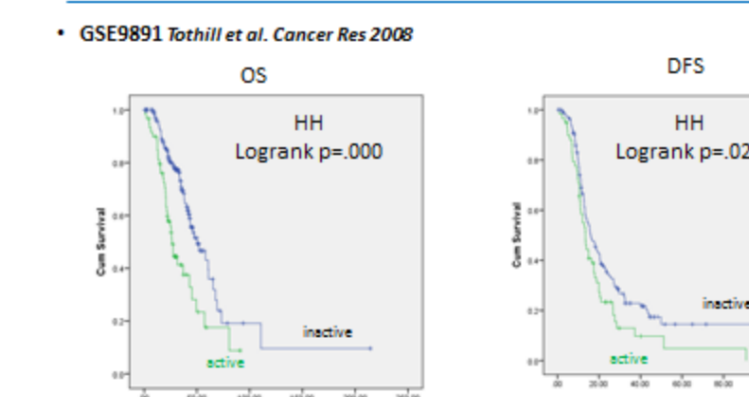
Histology	1	Serous HGS
FIGO stage	1	primary debulking surgery
IDS	1	interval debulking surgery
Recurrence	1	primary untreated disease (biopsy)
Chemotherapy	CT	Carboplatin/Paclitaxel
FIGO	CisPT	Cisplatin/Paclitaxel
PD	PD	progressive disease
nk	nk	not known

#### Public (GEO) datasets HGS

array	sample	FOXO	HH	NFkB	PI3K inact	PI3K act	HH act
GSM797484	516	7.7	-4.0	18.6	X		
GSM797485	518	6.9	-9.5	8.7	X		
GSM797486	519	-6.6	-4.2	2.7			
GSM797487	520	8.7	-1.9	15.9	X		
GSM797488	521	9.0	-7.1	11.0	X		
GSM797489	522	-0.2	-1.6	10.8			
GSM797490	523	-4.2	8.9	-3.6		X	
GSM797491	525	-1.9	0.2	12.9		X	
GSM797492	526	17.4	-5.4	25.4	X		
GSM797493	527	20.7	-6.0	23.8	X		

GSE32062: Advanced HGS ovarian cancer: HH/PI3K pathway combination

#### Public dataset Results - only HGS ovarian cancer samples: PFS

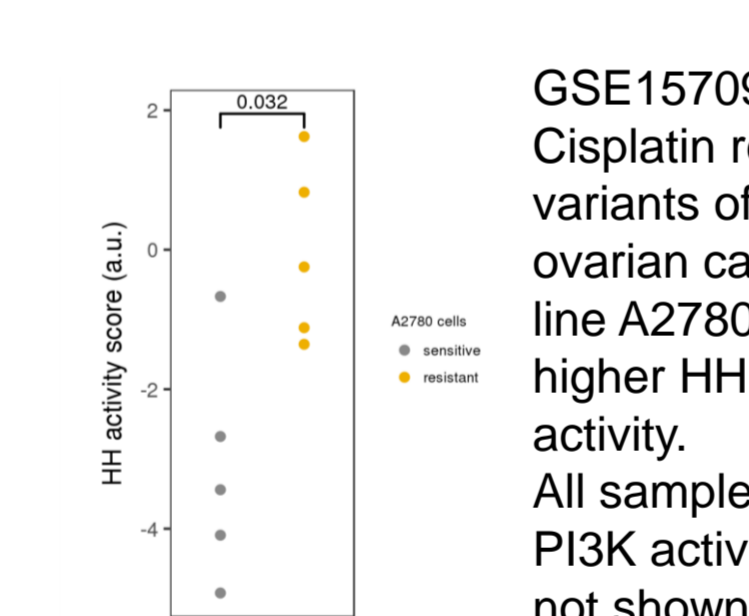


HH pathway activity → bad prognosis

#### Ovarian cancer cell lines

array	sample	FOXO	HH	NFkB
GSM341221	Carboplatin 24 hours set 1	-1.7	-4.6	3.5
GSM341222	Carboplatin 24 hours set 2	-2.0	-4.8	3.6
GSM341223	Carboplatin 30 hours set 1	-0.2	-5.1	6.8
GSM341224	Carboplatin 30 hours set 2	1.2	-4.6	7.7
GSM341225	Carboplatin 36 hours set 1	0.2	-5.1	8.8
GSM341226	Carboplatin 36 hours set 2	1.4	-6.4	9.0
GSM341227	Vehicle-Control 24 hours set 1	-1.6	-3.2	3.1
GSM341228	Vehicle-Control 24 hours set 2	-4.8	-3.4	0.8
GSM341229	Vehicle-Control 30 hours set 1	-3.4	-4.4	2.0
GSM341230	Vehicle-Control 30 hours set 2	-4.3	-5.2	1.2
GSM341231	Vehicle-Control 36 hours set 1	-1.4	-5.2	2.9
GSM341232	Vehicle-Control 36 hours set 2	-0.2	-4.5	2.3

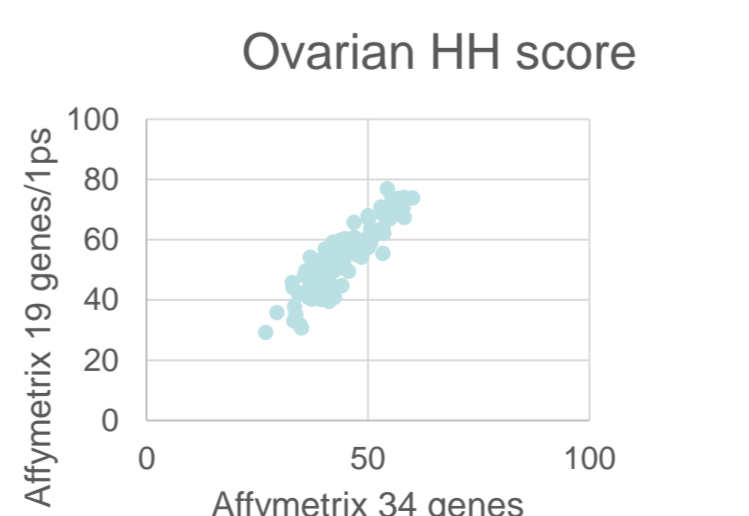
GSE15709: Cisplatin resistant variants of HGS ovarian cancer cell line A2780 have higher HH pathway activity. All samples are PI3K active (data not shown)



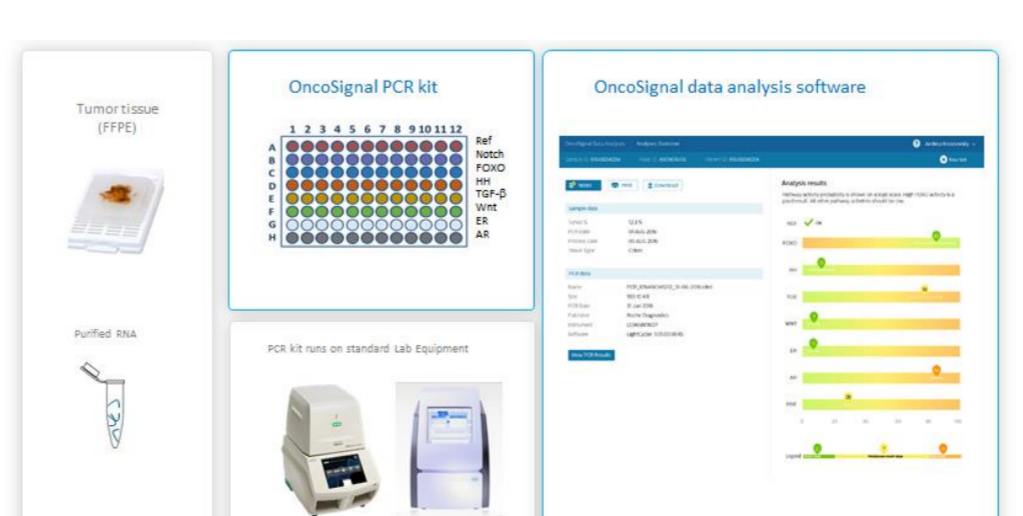
### Translation to qPCR for routine diagnostics

The Affymetrix models were ported to an RT-qPCR platform with 96 real-time qPCR assays.

Translation from 34 target genes to 12 genes for qPCR plate



OncoSignal qPCR test procedure and cloud-based pathway activity analysis



### Conclusion

Functional signal transduction pathway activity can be quantitatively measured using Philips pathway analysis on Affymetrix microarray and RT-qPCR data. Combined patient studies (including public GEO datasets) and cell line results suggest that HH/PI3K pathway activity in HGS ovarian cancer is associated with bad prognosis and platin resistance.

### Summary results

- HGS: two subtypes: HH/PI3K pathway active and FOXO/NFkB pathway active.
- HH activity associated with bad prognosis
- Patient with high HH/PI3K pathway activity: primary platin resistance and rapid progression.
- Ovarian cancer cell line (A2780): HH/PI3K pathway activity associated with platin resistance.
- Activity of NFkB/FOXO may be associated with less aggressive phenotype. In platin-sensitive ovarian cancer cell line, platin induces NFkB/FOXO activity, associated with apoptotic changes.

### Discussion

HH activity causing platin resistance has been described for gastric cancer (Yoon et al, Clin Cancer Res. 2014;20(15):3974-88.) Possible mechanism: HH transcription factor GLI induces expression of the Nucleotide Excision Repair protein XPD, which can repair platin-induced DNA damage. Signal transduction pathway analysis may provide valuable information to decide on targeted therapy and immune therapy choice.