

Fibroblast Growth Factors: New Drugs to Manage a Morpheic Tumor Target

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Disclosure

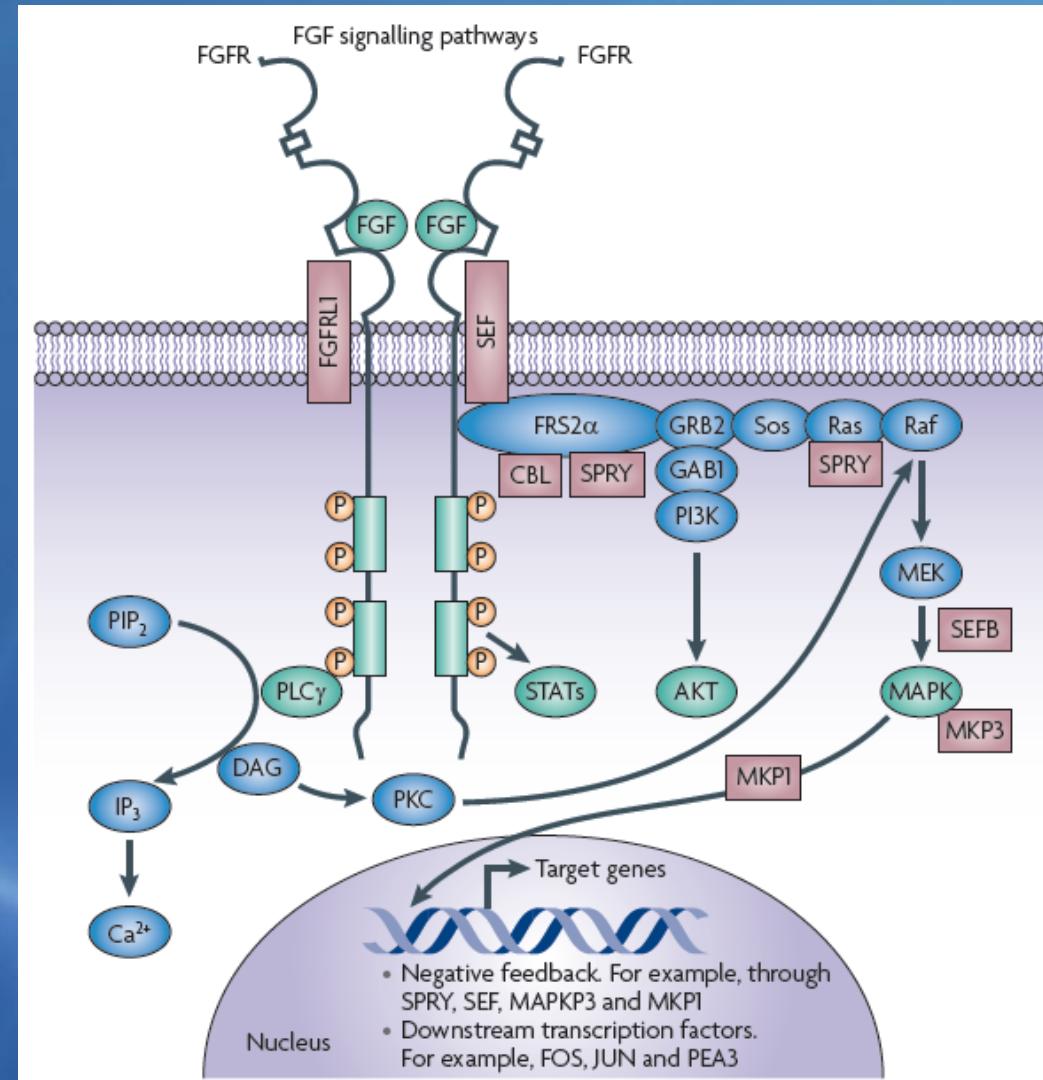
Consultant : Bristol-Mayers ; Novartis ;
Roche; Boehringer ; GSK ; Eisai ; Philogen

Angler :



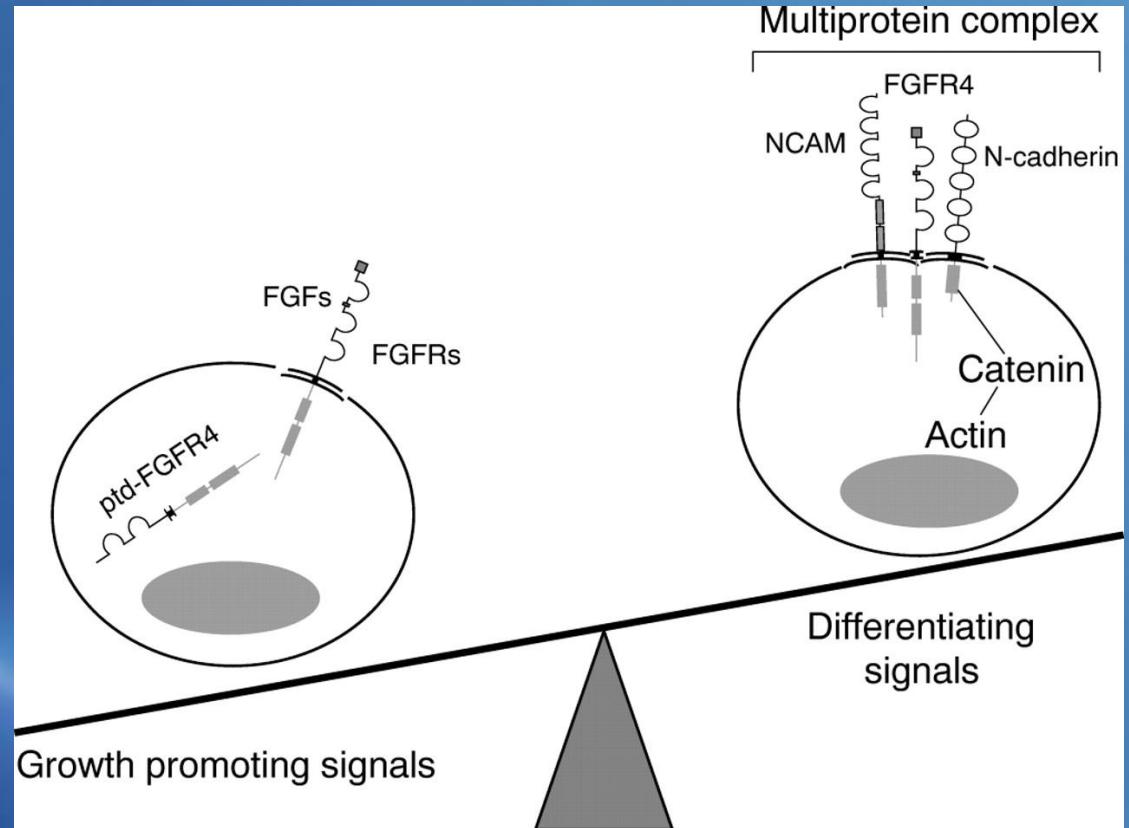
FGF/FGFR network

- 18 ligands
- 23 “members”
- 4 TK receptors: FGFR1, FGFR2, FGFR3, FGFR4
- 1 non TK receptor: FGFLR1 (negative control)
- Regulatory activity of cell proliferation, survival, migration and angiogenesis



Context-dependent activity

- In Normal Conditions
- In Cancer
 - Tumor suppressive
(
Medulloblastoma,
Prostate)
 - Tumor Driver



Ezzat et al. J Clin Pathol 2005

Microenvironmental Control To Be Clarified

FGF signalling in Tumors

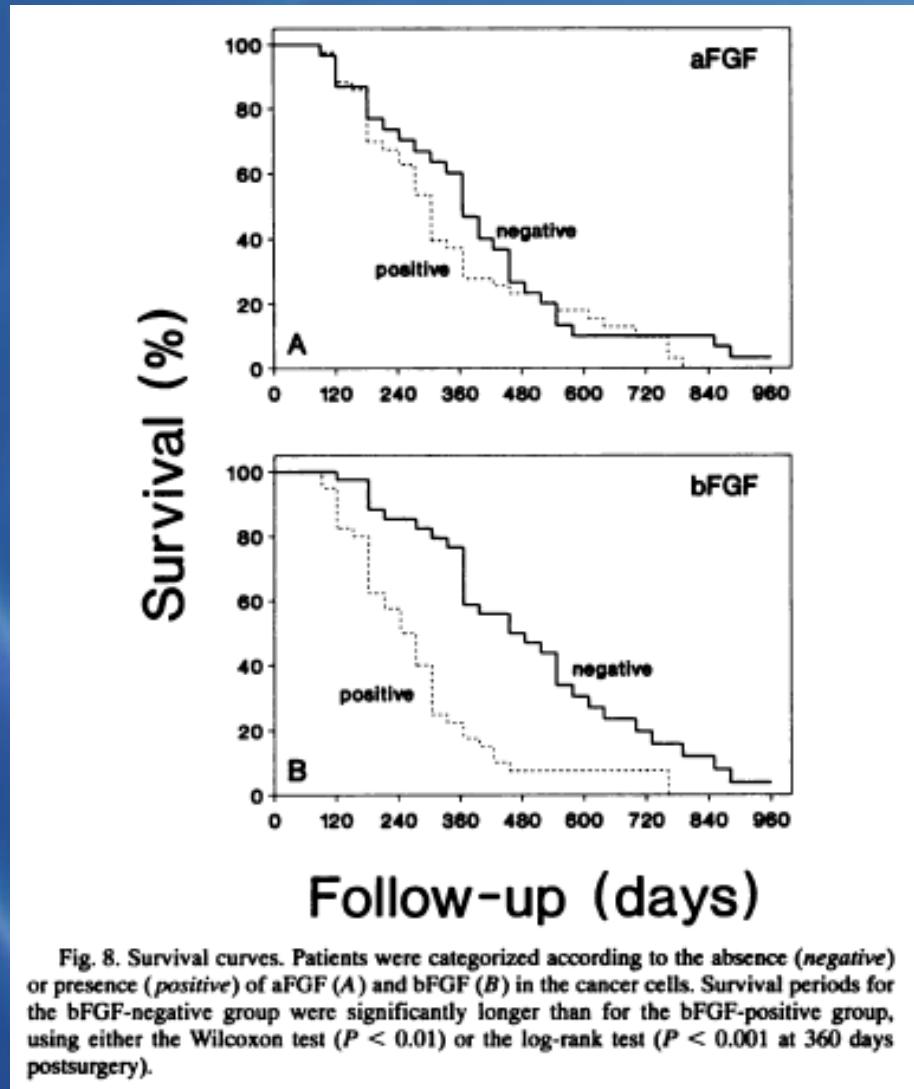


Fig. 8. Survival curves. Patients were categorized according to the absence (*negative*) or presence (*positive*) of aFGF (A) and bFGF (B) in the cancer cells. Survival periods for the bFGF-negative group were significantly longer than for the bFGF-positive group, using either the Wilcoxon test ($P < 0.01$) or the log-rank test ($P < 0.001$ at 360 days postsurgery).

Deregulation of FGF Signalling in Cancer

FGFR (4 + 1)

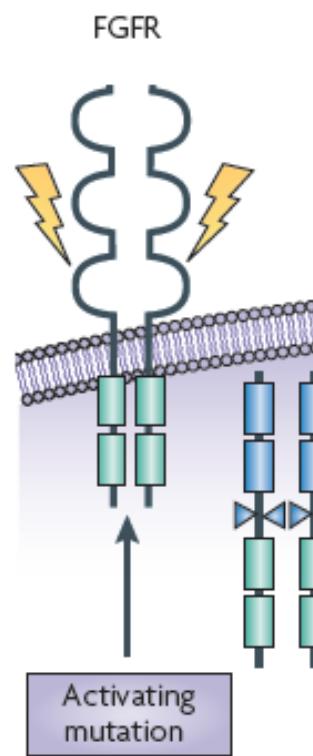
- ACTIVATING MUTATIONS
- GENE AMPLIFICATION
- CHROMOSOMAL TRASLOCATIONS

FGF (18 + 5)

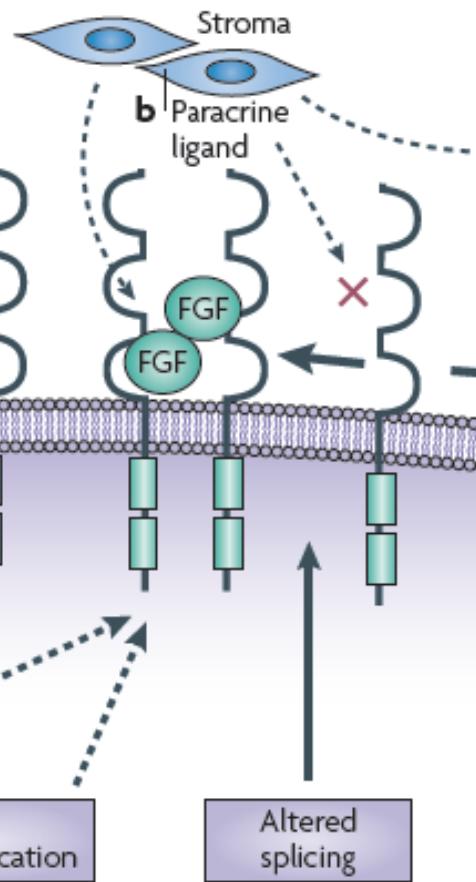
- AUTOCRINE SIGNALLING (ie. Melanoma)
- PARACRINE SIGNALLING (ie. Prostate cancer)
- HORMONAL SIGNALLING (ie. HCC)

Mechanisms of pathogenetic cancer cell FGF signalling

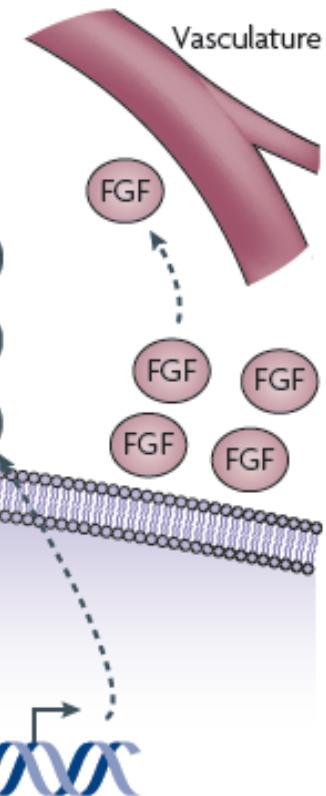
a Ligand-independent signalling



Ligand-dependent signalling



d Angiogenesis



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Turner et al. Nat Canc Rev 2010

Tumor-specific genetic FGFR alterations

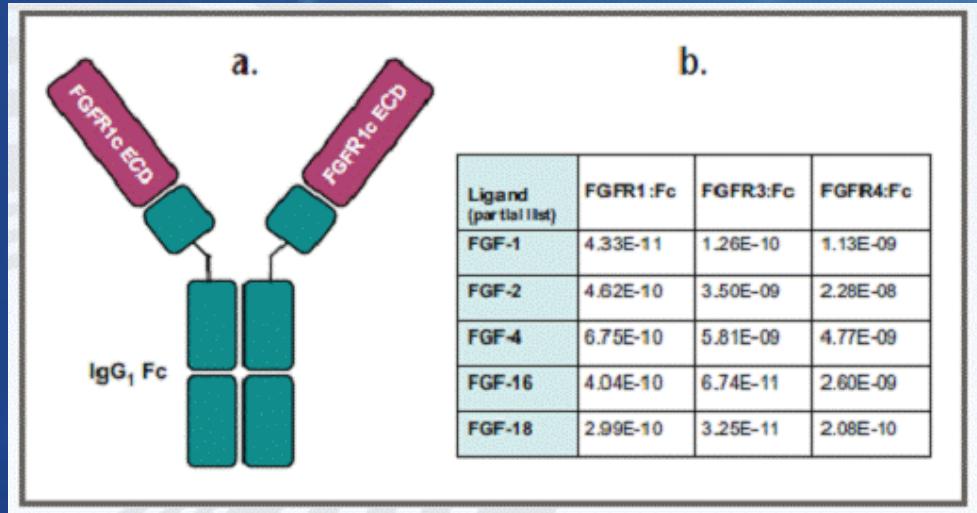
| FGFR gene | Cancer type | Germline or somatic | Genetic changes |
|--|-----------------------------|---------------------|---------------------------|
| FGFR1 | Breast cancer | Somatic | Gene amplification |
| | Ovarian cancer | Somatic | Gene amplification |
| | Myeloproliferative syndrome | Somatic | Chromosomal translocation |
| | Bladder cancer | Somatic | Gene amplification |
| | Glioblastoma | Somatic | Missense mutation |
| | Oral cancer | Somatic | Gene amplification |
| | Melanoma | Somatic | Missense mutation |
| | Rhabdomyosarcoma | Somatic | Gene amplification |
| FGFR2 | Breast cancer | Somatic | Gene amplification |
| | | Somatic | Missense mutation |
| | | Germline | Intronic regulatory SNPs |
| | Gastric cancer | Somatic | Gene amplification |
| | | Somatic | Missense mutation |
| | Lung cancer | Somatic | Missense mutation |
| | Uterus cancer | Somatic | Missense mutation |
| | Melanoma | Somatic | Missense mutation |
| FGFR3 | Multiple myeloma | Somatic | Chromosomal translocation |
| | | Somatic | Missense mutation |
| | Bladder cancer | Somatic | Missense mutation |
| | | | Gene amplification |
| | Cervical cancer | Somatic | Missense mutation |
| | Colorectal cancer | Somatic | Missense mutation |
| | Peripheral T-cell lymphoma | Somatic | Chromosomal translocation |
| | Oral cancer | Somatic | Missense mutation |
| FGFR4 | Testicular tumor | Somatic | Missense mutation |
| | Breast cancer | Germline | Missense coding SNP |
| | Lung cancer | Germline | Missense coding SNP |
| | Rhabdomyosarcoma | Somatic | Missense mutation |
| FGFR: FGF receptor; SNP: Single-nucleotide polymorphism. | | | |



Anti FGFR/FGFs Monoclonal Antibodies

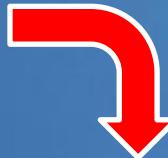
| Drug (company) | Target | Clinical Development |
|---------------------------|-----------------|---|
| FP-1039 (Five Prime Ther) | FGF ligand trap | Phase I moving to phase II in Endometrial Cancer |
| Anti-FGFR3 (Genentech) | FGFR3 | Phase I Multiple Myeloma |
| IMC-A1 (ImClone) | FGFR1 | NA |
| PRO-001 (ProChon Biotech) | FGFR3 | NA |
| 1 A6 (Genentech) | FGF19 | NA |

FP-1039



1) **Structure:** extracellular domains of human FGFR1 isoform α-IIIc + native Fc regions of human IgG₁.

2) **Single agent:** growth inhibition of 4/12 pts-derived breast cancer xenografts and vs. H460 (lung), Colo205 (colon) and Caki-1 (renal) models., **Combo:** + sunitinib or + CDDP or + bevacizumab resulted in additive anti-tumor activity in H460 and Colo205 xenografts.



2 Clinical Trial Ongoing:

1) Safety Study of FP-1039 To Treat Cancer

2) Study of FP-1039 in Subjects With Endometrial Cancers



Pts selection according to the presence of the point mutation S252W in the extra domain of FGFR2 (~7% of all cases)

Long et al. Proc Am Assoc Cancer Res; 2009; abstract 2789.

Keer et al, J Clin Oncol 28:15s, 2010; abstr TPS260).
www.clinicaltrial.gov

FGFR TKI

| Agent | Company | Target | Clinical Development |
|--------------------|-------------------------|---|---|
| Brivanib alaninate | BMS | VEGFR 1-2-3 FGFR 1-2-3 | Phase III |
| Masitinib | AB Science | FGFR3, c-kit, PDGFR, FAK | Phase III (pancreas, GIST, multiple myeloma) |
| BIBF1120 | Boehringer Ingelheim | VEGFR 1-2-3 FGFR 1-2-3 PDGFR a & B | Phase III |
| TKI258 (Dovitinib) | Novartis | VEGFR 1-2-3, FLT, FGFR 3 cKit , PDGFR B | Phase II |
| TSU-68 | Taiho Pharm. | VEGFR1, FGFR 1 PDGFR B | Phase I/II |
| E-3810 | EOS | VEGFR ; FGFR1 | Phase I |
| E7080 | Eisai | FGFR, PDGFR, VEGFR | Phase I |

AP24534 (Panatinib) IC₅₀ of 0.37, 2, 1.5, 2.2, 1.1, 1 and 0.24 nM for native pan-BCR-ABL, mutated form, VEGFR2, FGFR1, PDGFR α , mutant FLT3 and LYN.

BIBIF1120 (Vargatef) all three VEGFR subtypes (IC₅₀=13-34nmol/L) , PDGFR α and PDGFR β (IC₅₀=59and65nmol/L) , FGFR 1, 2, and 3 (IC₅₀=69, 37, and 108 nmol/L)

BMS -582664 (Brivanib alaninate) VEGFR-2(25 nM), VEGFR-1 (380nM), VEGFR-3 (10 nM); FGFR 1 , 2 , 3 (IC₅₀=148 nM, 125 nM, 68 nM)

TKI-258(Dovitinib) IC₅₀ of 1, 2, 5, 10, 8, 27, 36 nM for FLT3, c-KIT, FGFR3, VEGFR1/2/3, PDGFR β and CSF-1R, respectively

TSU-68 (SU6668) VEGF-R1, PDGF-R β and FGF-R1 (IC₅₀ : 2.1 μ M, 8 nM and 1.2 μ M)

Ki8751 selective VEGFR-2 tyrosine kinase inhibitor with IC₅₀ of 0.9 nM, 40, 67, 170 nM for VEGF-2, c-Kit, PDGFR α and FGFR-2, respectively

PD 173074 Inhibitor of FGFR1 (IC₅₀ = 21.5 nM) and VEGFR

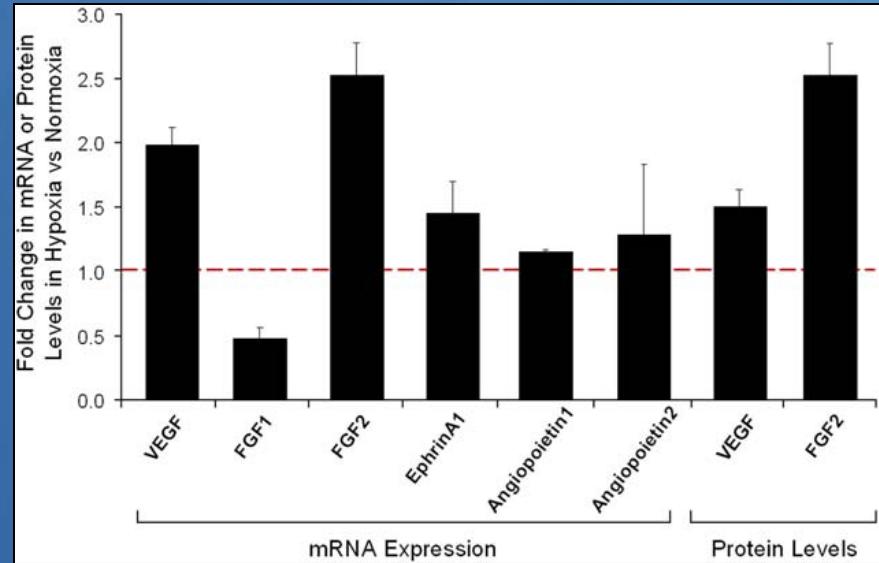
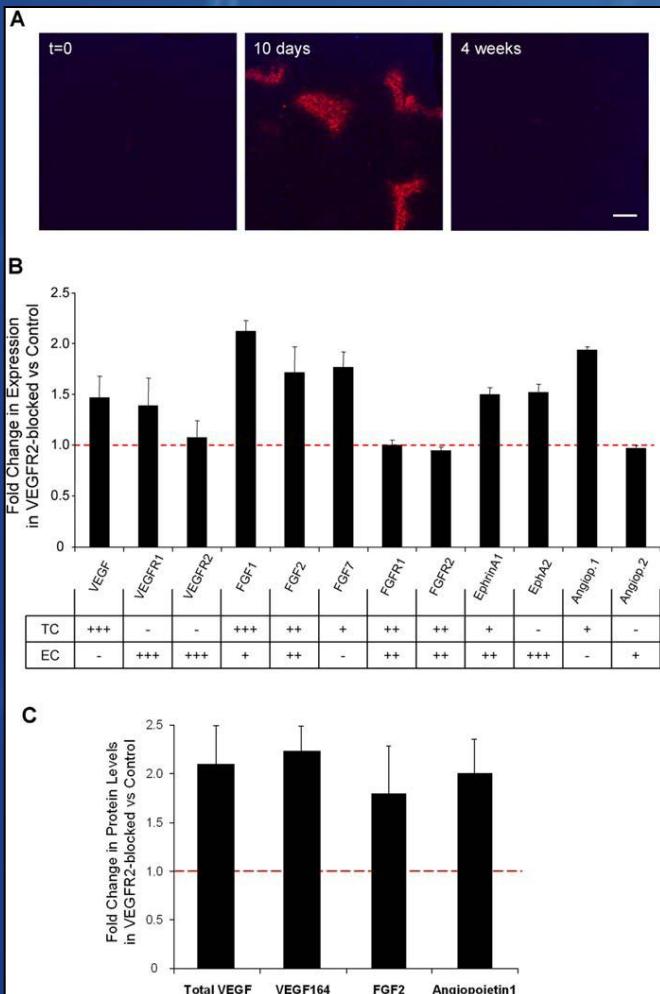
E-3810 VGFR 1,2,3 (IC₅₀ : 7,25, 10 nM) FGRF 1,2 3 (IC₅₀ : 17.5, 81.5, 237.5 nM)

FGFR TK inhibitors

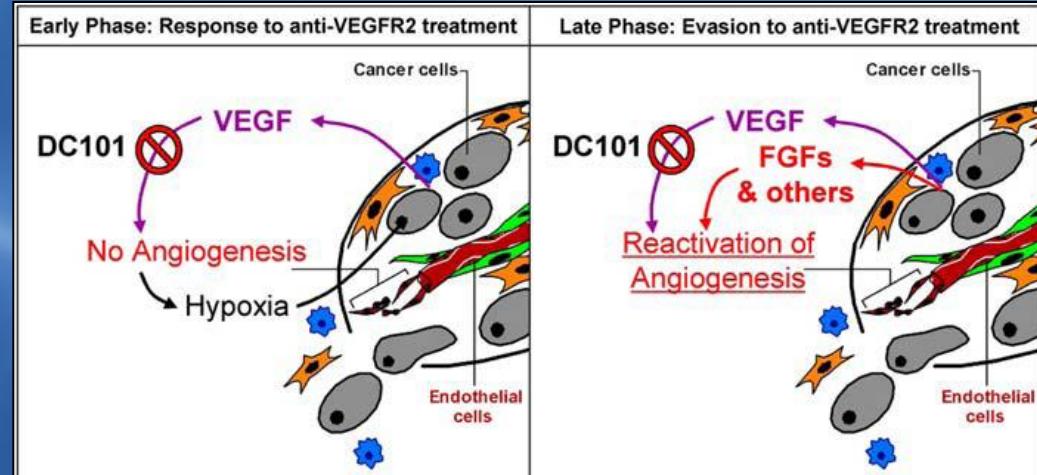
| | Sorafenib | Sutent | Brivanib | BIBF 1120 | Dovitinib | E-3810 |
|--------|-----------|--------|----------|-----------|-----------|---------|
| VEGFR1 | - | 2 | 350 | 34 | 10 | 7 |
| VEGFR2 | 90 | 9 | 26 | 21 | 8 | 25 |
| VEGFR3 | 20 | 17 | 10 | 13 | 27 | 10 |
| FGFR1 | - | - | 150 | 69 | Low | 17.5 |
| FGFR2 | 580 | 830 | 125 | 37 | - | 82.5 |
| FGFR3 | - | - | 68 | 108 | 5 | 237.5 |
| PDGFR | 57 | 8 | 7,460 | 59/60 | 36 | 175/525 |
| C-Kit | 68 | 13 | Low | - | 2 | 456 |
| Flt-3 | 58 | 10 | 58 | - | 1 | - |
| Other | RET,RAF | RET | RET | Src | CSF-1R | CSF1R |

Evasion of antiangiogenic targeting of VEGF

1. VEGFR2 blockade → hypoxia → change of expression of proangiogenic factors in tumors

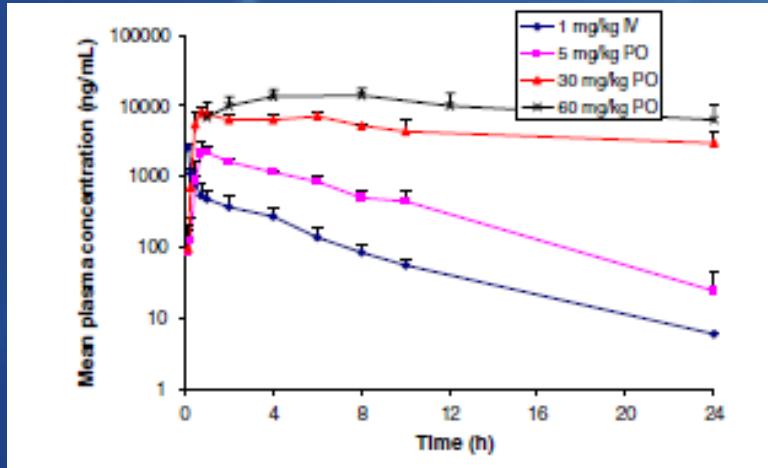


2. Hypoxia → ↑expression of proangiogenic factors in tumor-derived βTC cell line

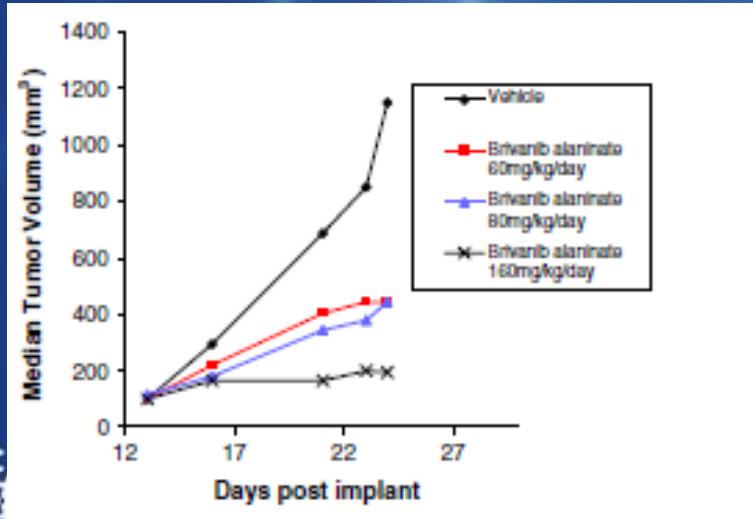


Brivanib (BMS-582664)

Preclinic: Plasma Concentrations



Preclinic: Antitumor Activity on HCT116/VM46 colon cancer q1Dx12



Clinic:

Prodrug of BMS-540215; dual inhibitor of VEGFR and FGFR

Linear PK

- No impact of race, age, gender
- No impact of mild liver impairment (Child-Pugh A)

800 mg QD PO continuously; no food effect

Eliminated mainly through metabolism

Weak Inhibitor of UGT1A1; no interaction with CYP3A4 substrates.

Jonker DJ et al. JCO, 2007: abstr. 3559
Marathe et al, Cancer Chemother Pharma 2009

Brivanib Single Agent Phase I Results

AEs from Phase I + Phase II (HCC)

| Side Effects (grade I-III) | Frequency |
|-------------------------------|-----------|
| Fatigue | 33-45% |
| Anorexia | 27-39% |
| Diarrohea | 14-35% |
| Nausea | 14-65% |
| Rash | 6-24% |
| Hypertonia | 9-28% |
| Emesis | 9-41% |

Phase I Antitumor Activity

| Response ^a | All Doses, n (%) |
|-----------------------|--------------------|
| Partial Response | 2 (3) ^b |
| Stable Disease | 24 (35) |
| Progressive Disease | 32 (47) |
| Not assessable | 10 (15) |

^a At least 1 post treatment CT;
^b 1 Renal cell carcinoma 600 mg q1D; 1 Ampulla of Vater 1000 mg q1D

Dempke et al. Anticancer Res 2010

Jonker DJ et al. Annals Oncol 2010

Phase II Brivanib Single Agent in HCC

Dose: Brivanib 800 mg PO QD

Cohort A: no previous systemic CT for HCC

Cohort B: 1 prior regimen with antivascular therapies

End Points: PFS, TTP, OS, Disease control rate, Safety.

| Baseline Characteristics | Cohort A (n=55) | Cohort B (n=46) |
|------------------------------------|-----------------|-----------------|
| Age | 60 (27-80) | 55 (21-81) |
| Male/Female, % | 89/11 | 72/28 |
| Asian/non-Asian, % | 64/36 | 67/33 |
| HBV+/HCV-, % | 53/22 | 65/17 |
| ECOG PS 0/1/2, % | 45/49/5 | 26/72/2 |
| Extrahepatic spread, % | 76 | 78 |
| Child-Pugh A/B, % | 91/9 | 91/9 |
| AFP >UNL, % | 76 | 83 |
| Local Therapy (TACE, RFTA etc.), % | 49 | 83 |

Brivanib induces responses in HCC pts



04 Jul 07

Baseline
assessment



18 Mar 08

mWHO: SD
mRECIST: CR



12 May 09

mWHO: CR
mRECIST: CR

Park JW et al. Clin Cancer Res 2011



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Tumor responses: mWHO and mRECIST

| Outcome | mWHO | mRECIST(3, 22) |
|---|------------------------------|------------------|
| Overall Survival (95% CI) (median, months) | 10 (95% CI, 6.8-15.2) | |
| Best tumor response, n (%) | | |
| CR | 1 | 5 (9.1) |
| PR | 3 (5.5)) | 9 (16.4) |
| SD | 24(43.6) | 29 (52.7) |
| SD | 22 (40.0) | |
| uCR | 0 | 2 (3.6) |
| uPR | 2 (3.6) | 3 (5.5) |
| DCR (CR + PR + SD), n/N (%) | 28/55 (50.9) | 43/55 (78.2) |
| Median TTP(95% CI) (months) | 2.8 (1.4–3.5) ^a | 5.4 (2.8,–) |
| Median PFS (months) | 2.7 (1.4-3.0) ^a | 4.7 (2.8-5.7) |
| 6-month PFS rate (%) | 21.1 (9.6-32.5) ^a | 35.6 (21.0-49.4) |

^aIRRC assessment



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Park JW et al. Clin Cancer Res 2011

BIBF 1120

1) Linear PK

2) MTD: 250 mg PO QD and BID

3) RD: 200 mg PO BID

4) AEs grade 1-2: nausea (68.9%), vomiting (45.9%), diarrhoea (44.3%).

5) Responses: 2 PR (RCC, CRC), 1 CR (RCC)

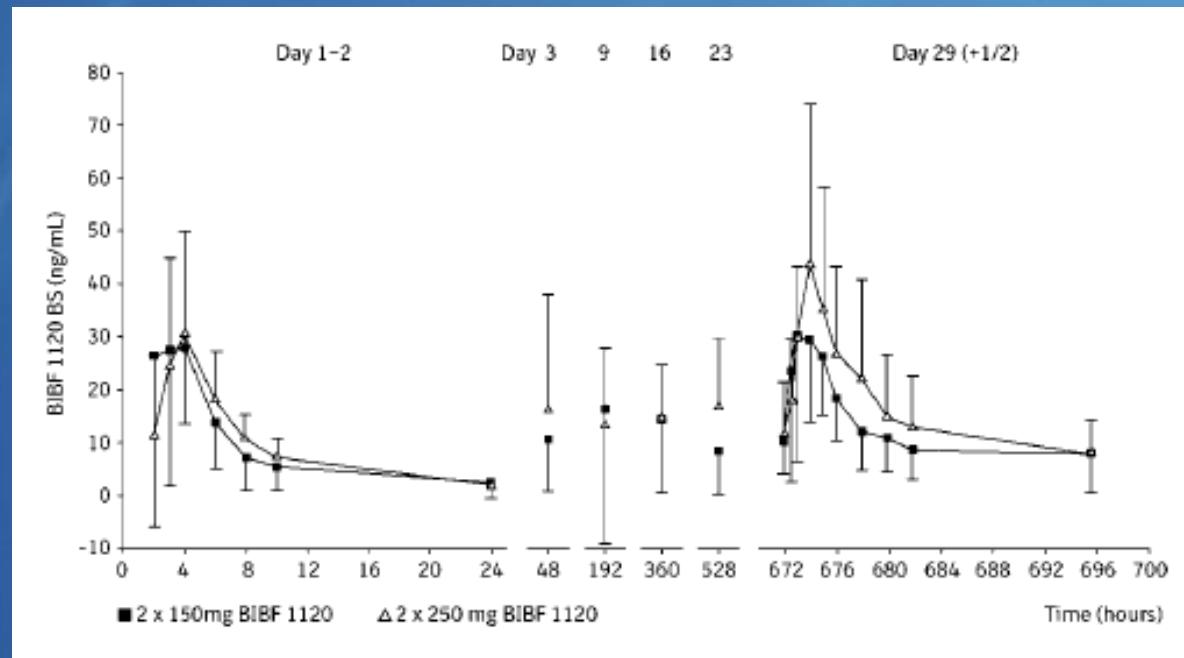


Table 3. Number of patients with DLT within the first course of BIBF 1120 treatment

| Total daily dose (mg) | Once daily | | Twice daily | |
|-----------------------|--------------|---------------------------------------|--------------|---|
| | All patients | Patients with DLT | All patients | Patients with DLT |
| 50 | 2 | 0 | — | — |
| 100 | 1 | 0 | — | — |
| 150 | — | — | 6 | 0 |
| 150 + 200 | — | — | 6 | 0 |
| 200 | 8 | 1 (12.5) AST+γGT increased | 6 | 1 (16.7) CD4 lymphocytes decreased |
| 250 | 6 | 1 (16.7) CD4 lymphocytes decreased | 13 | 1 (7.7) nausea |
| 300 | 5 | 2 (40) AST+γGT (1), ALT+AST (1) | 5 | 4 (80) nausea+vomiting (1), ALT+AST increased (1), ALT+AST+γGT (1), CD4 lymphocytes decreased + γGT increased (1) |
| 450 | 3 | 2 (66.7) ALT+AST+γGT (1), ALT+γGT (1) | — | — |

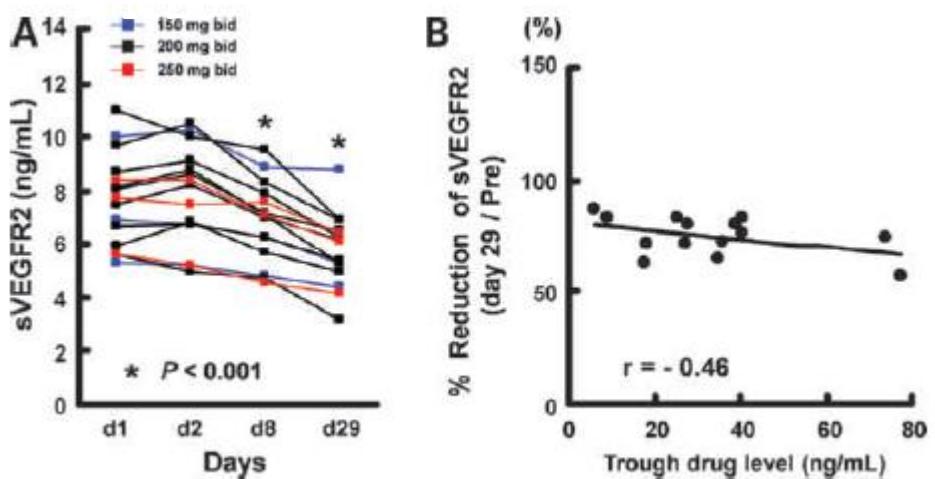
NOTE: DLT was defined as any CTC grade 3 or 4 hematologic or nonhematologic toxicity.

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Mross K et al. Clin Cancer Res 2010

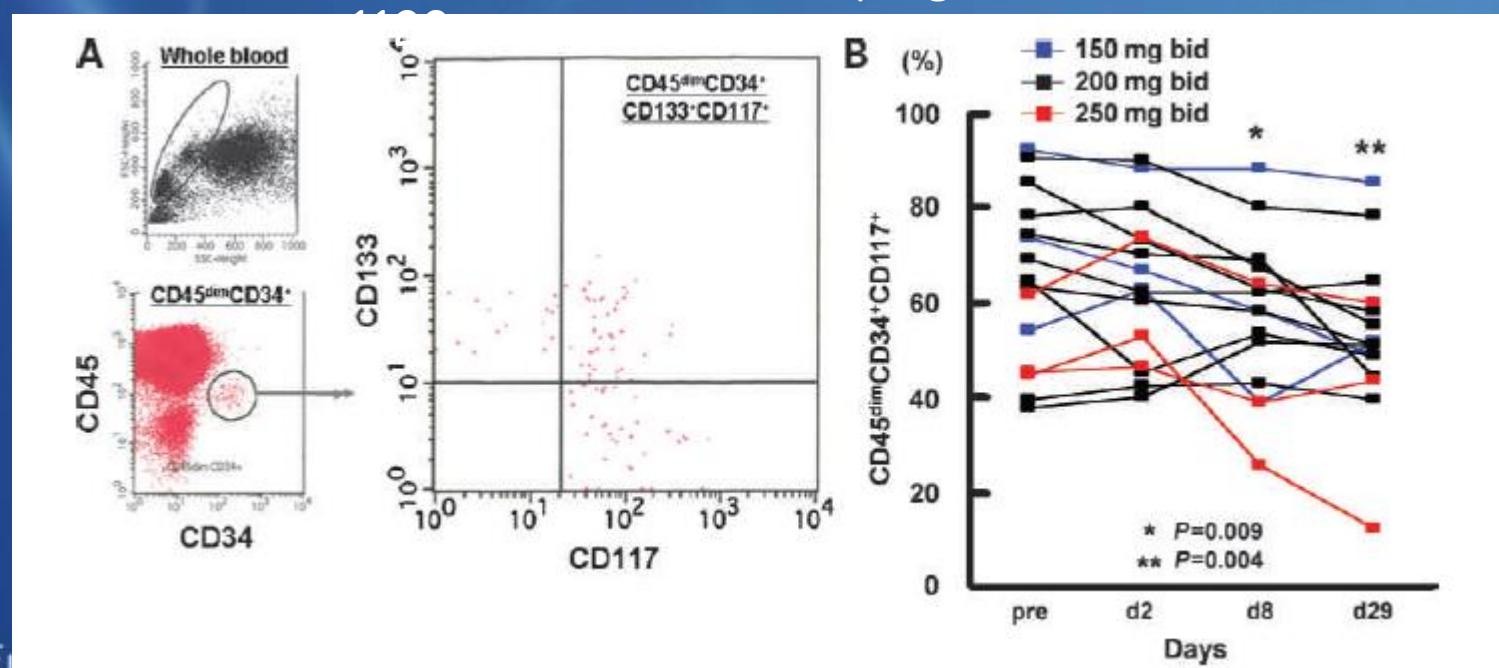
1) VEGF values after BIBF 1120



Biomarkers study of BIBF 1120

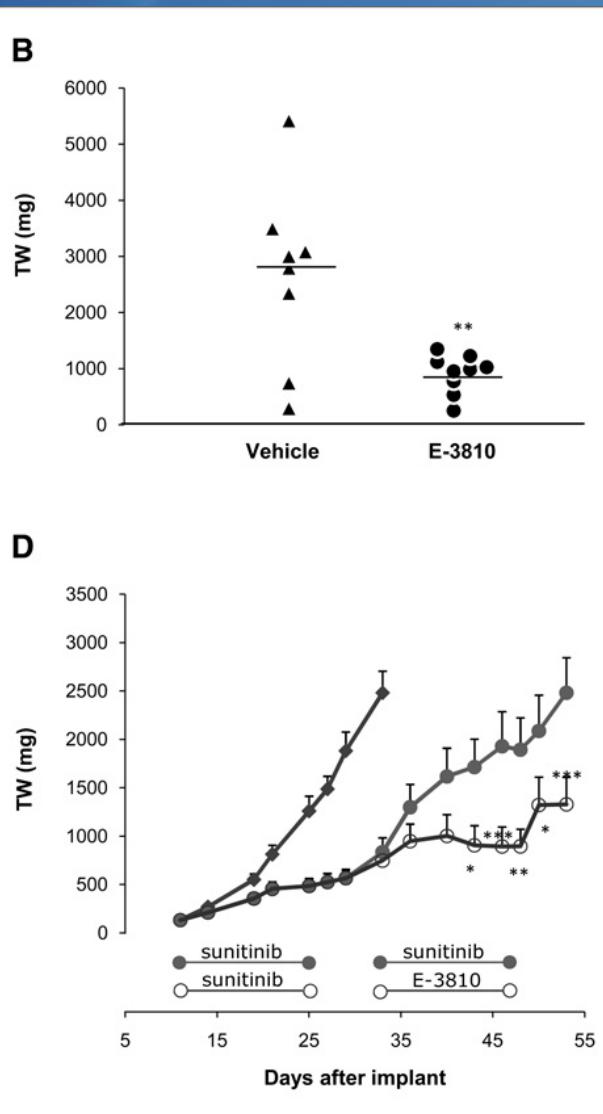
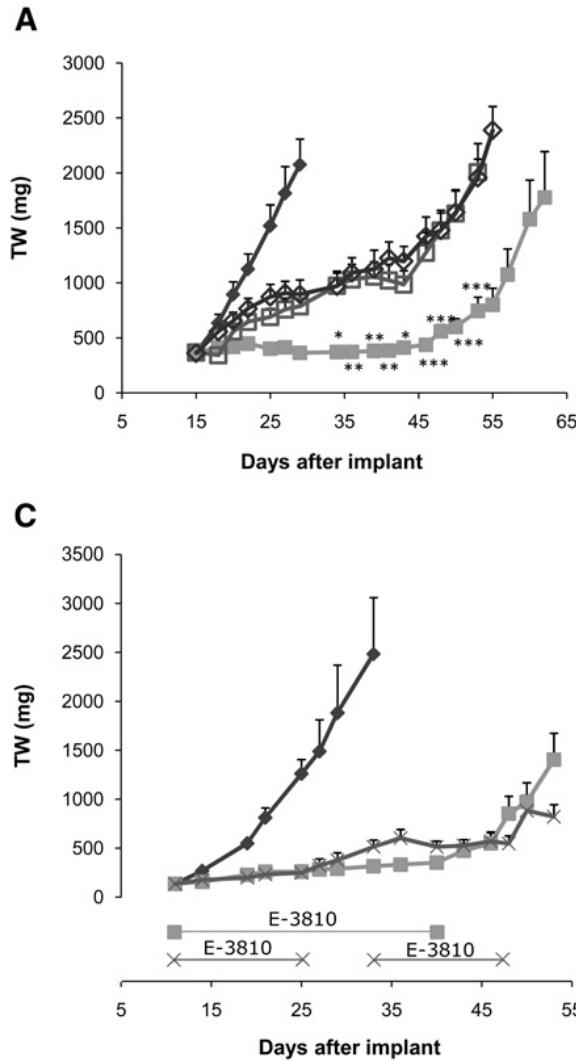
Okamoto et al. Mol Cancer Ther 2010

2) Serum CD117 BMD progenitor cell after BIBF 1120



E-3810

Antitumor activity



Dual VEGFR and FGFR1 inhibitor

Phase I trial single agent ongoing



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Bello et al. Cancer Res 2011

Future directions: Combo with CT

Phase I Trial of BIBF 1120 + Pemetrexed in NSCLC

Pts population: 26 pts with NSCLC relapsed after 1 previous Cisplatin-based 1st line for advanced disease

Doses:

Pemetrexed 500 mg/m² (day 1 of every cycle)

BIBF 1120: PO BID 100, 150, 200, 250 mg (from day 2 of every cycle)

RD: BIBF 1120 → 200 mg PO BID

Antitumor activity: 1 CR, 13 SD, 8 PD, 3 missed FUP, 1 non evaluable

| | All CTCAE grades, n (%) | CTCAE grade 3,* n (%) |
|----------------------|-------------------------|-----------------------|
| Fatigue | 17 (65.4) | 6 (23.1) |
| Nausea | 16 (61.5) | 1 (3.8) |
| Anorexia | 14 (53.8) | 2 (7.7) |
| Rash | 10 (38.5) | 0 |
| Diarrhea | 9 (34.6) | 1 (3.8) |
| Vomiting | 9 (34.6) | 1 (3.8) |
| ALT increases | 7 (26.9) | 3 (11.5) |
| Abdominal pain | 6 (23.1) | 2 (7.7) |
| Dysgeusia | 6 (23.1) | 0 |
| Pruritus | 6 (23.1) | 0 |
| Insomnia | 5 (19.2) | 1 (3.8) |
| AST increases | 5 (19.2) | 0 |
| Dyspepsia | 4 (15.4) | 0 |
| Headache | 4 (15.4) | 0 |
| Constipation | 3 (11.5) | 0 |
| Stomatitis | 3 (11.5) | 0 |
| Chills | 3 (11.5) | 0 |
| Dermatitis acneiform | 3 (11.5) | 0 |

NOTE: Data presented are the highest ever reached CTCAE grade. One patient may have experienced more than one event.

*No grade 4 adverse events were observed.



Future directions: Combo with Biologics

Phase I Trial of Brivanib + Cetuximab in GI tumors

Pts population: 60 pts with GI tumors relapsed after previous CTs (expansion cohort for CRC)

Doses:

Brivanib PO QD 250 → 800 mg D1-D8

Cetuximab 400 mg/m² C1D8 → 250 mg/m² qwk

G3/4 Aes: fatigue 16%, transaminitis 12%, diarrhoea 2%; 1 DLT: bilateral pulmonary emboli.

| | Prior Therapy | | | | |
|-----------------------|---------------|----|---|---|----------|
| EGFR inhibitor | - | - | + | + | |
| VEGFR inhibitor | - | + | - | + | |
| Evaluable pts (total) | 12 | 12 | 1 | 9 | 34 |
| PR | 4 | 1 | 0 | 0 | 5 (15%) |
| SD | 5 | 7 | 1 | 6 | 19 (56%) |
| PD | 3 | 4 | 0 | 3 | 10 (30%) |
| PR + SD ≥ 6 months | 4 | 3 | 0 | 3 | 10 (30%) |

Garrett CR et al. JCO 2008: abstr. 4111

Conclusions

- 1) FGFs/FGFRs signalling is a major driver of carcinogenesis. It is tumor specific
- 2) TK inhibitors not specific for FGFR signalling have a potential advantage on evasion of antiangiogenic treatments
- 3) Promising results have been obtained in single agents trials
- 4) Future directions require :
 - To improve knowledge to select patients and tumors
 - To explore combination or sequential schedules