

Tyrosine kinase inhibitor BIBF1120 ameliorates inflammation, angiogenesis and fibrosis in CCl₄-induced liver fibrogenesis mouse model

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Supplementary Table 1: Antibodies used for the immunohistochemistry

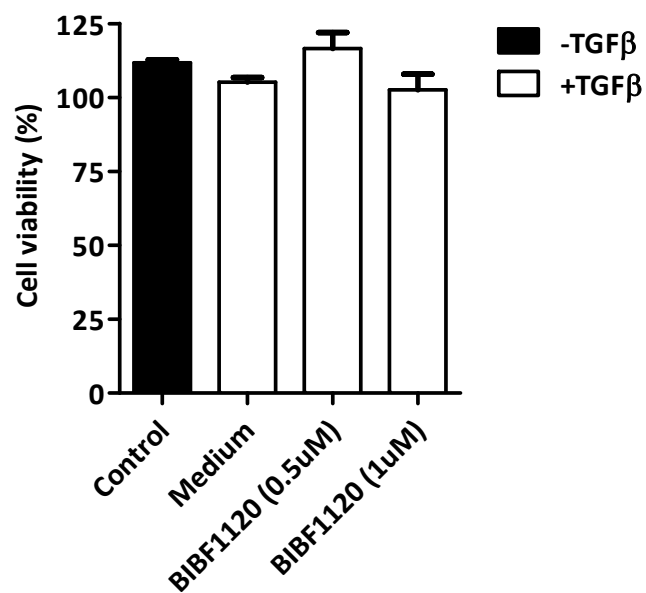
Primary Antibody	Source
Monoclonal mouse anti- α -SMA	Sigma
Polyclonal goat anti-collagen I	Southern Biotech
Monoclonal rat anti-mouse CD31	Southern Biotech
Polyclonal goat anti-desmin	Santa Cruz
Polyclonal goat anti-vimentin (sc-7557)	Santa Cruz
Monoclonal rat anti-MHC Class II: sc-59318	Santa Cruz
Polyclonal goat anti-YM-1 (Biotinylated Anti-mouse chitinase-3-like 3/ECF-L)	R and D systems
Monoclonal mouse anti- β -actin	Sigma
Polyclonal rabbit anti-ITGA5	Sigma
Polyclonal rabbit anti-phospho-FAK antibody	Cell signaling
Secondary Antibody	Source
Polyclonal goat anti-rabbit IgG	DAKO
Polyclonal rabbit anti-mouse IgG	DAKO
Polyclonal goat anti-mouse IgG	DAKO
Polyclonal rabbit anti-goat IgG	DAKO
Polyclonal goat anti-rat IgG	Southern Biotech

Supplementary Table 2: Sequence of the mouse primers used for quantitative real-time PCR

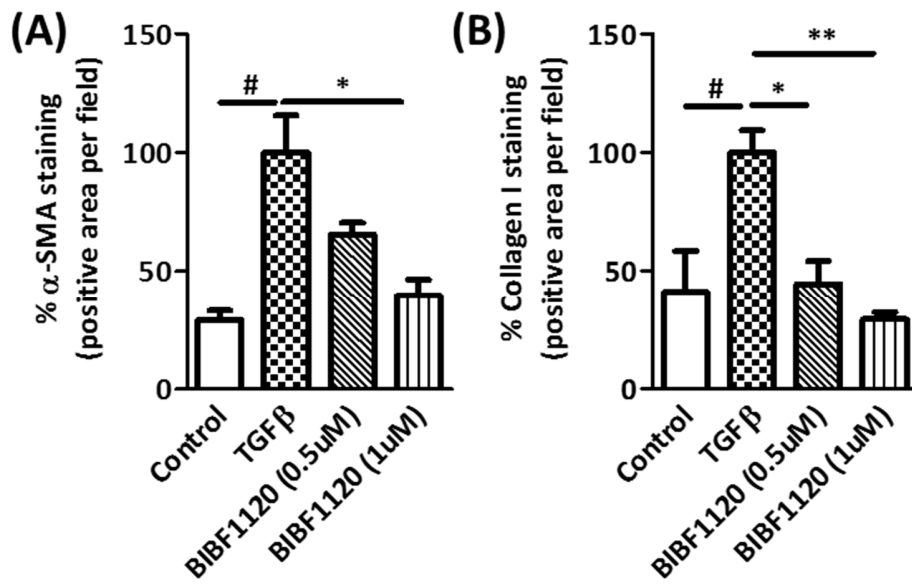
Gene	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
Collagen 1 α 1	TGACTGGAAGAGCGGAGAGT	ATCCATCGGTCATGCTCTCT
Desmin	ATGCAGCCACTCTAGCTCGT	CTCATACTGAGCCCGGATGT
ACTA2 (α -SMA)	ACTACTGCCGAGCGTGAGAT	CCAATGAAAGATGGCTGGAA
GAPDH	ACAGTCCATGCCATCACTGC	GATCCACGACGGACACATTG
TIMP1	ATCAGTGCCTGCAGCTTCTT	TGACGGCTCTGGTAGTCCTC
PDGF β R	GCTGGAGCTGAGTGAGAGTC	GCAGGTAGACCAGGTGACAT
FGFR1	GCTGACTCCAGTGCATCCAT	ACACGGTTGGGTTTGTCTT
VEGFR2	TAACCTGGCTGACCCGATTC	AAGTCACAGAGGCGGTATGC
CD31	TCCCTGGGAGGTCGTCCAT	GAACAAGGCAGCGGGGTTTA
CD34	GGGTAGCTCTCTGCCTGATG	TCTCTGAGATGGCTGGTGTG
Cadherin 11	CCGACTTGTGAATGGGACTCG	AGGGCCACAAAGCACAGTAA
SOX9	GTGCAAGCTGGCAAAGTTGA	TGCTCAGTTCACCGATGTCC
ITGA5	GAACCCTGTGTCCTGCATCA	TTGGAGTTCCACCTCGAAGC
Fibronectin	ATGAGAAGCCTGGATCCCCT	GGAAGGGTAACCAGTTGGGG
IL-6	TGATGCTGGTGACAACCACGGC	TAAGCCTCCGACTTGTGAAGTGGTA
TNF α	AGGCTGCCCGACTACGTGC	CAGCGCTGAGTTGGTCCCCC
CCL2 or MCP1	GTGCTGACCCCAAGAAGGAA	GTGCTGAAGACCTTAGGGCA
TIMP1	ATCAGTGCCTGCAGCTTCTT	TGACGGCTCTGGTAGTCCTC
Arg1	GTGAAGAACCCACGGTCTGT	CTGGTTGTCAGGGGAGTGTT
NOS2 or iNOS	AATCTTGGAGCGAGTTGTGG	CAGGAAGTAGGTGAGGGCTTG
Periostin	ATCCACGGAGAGCCAGTCAT	TGTTTCTCCACCTCCTGTGG
NOS3 or eNOS	CATGGGCAACTTGAAGAGTGT	GGGTGTCGTAGGTGATGCTG
IL-1 β	GCCAAGACAGGTCGCTCAGGG	CCCCCACAGTTGACAGCTAGG

Supplementary Table 3: Sequence of the human primers used for quantitative real-time PCR

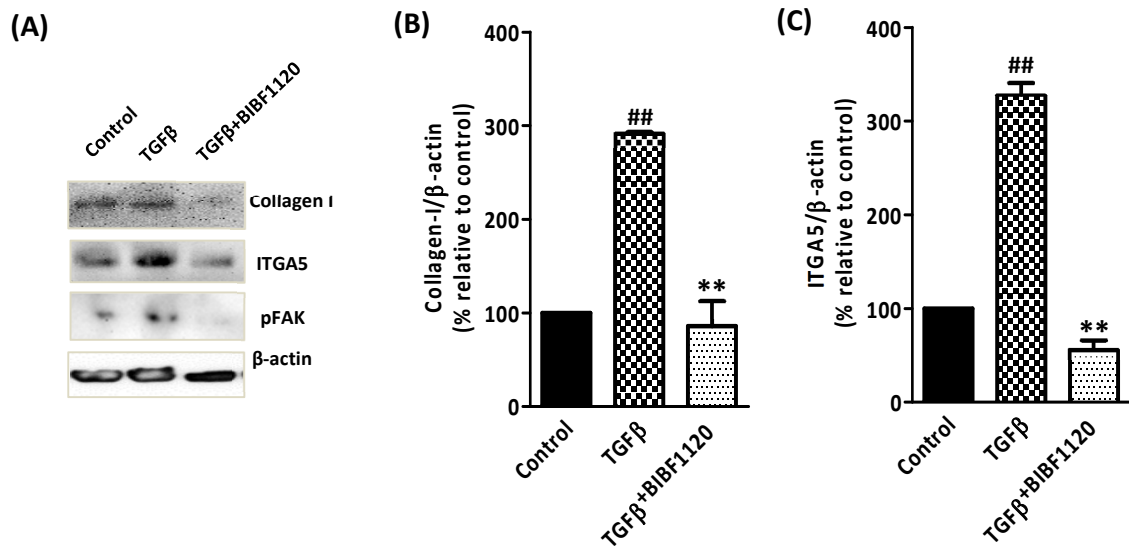
Gene	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
Collagen 1 α 1	GTACTGGATTGACCCCAACC	CGCCATACTCGAACTGGAAT
GAPDH	TCCAAAATCAAGTGGGGCGA	TGATGACCCTTTTGGCTCCC
PDGF β R	CATGGGGGTATGGTTTTGTC	GTAAGGTGCCAACCTGCAAT
TGF β 1	GCGTGCTAATGGTGAAACC	GAGCAACACGGGTTCAGGTA
TIMP1	GGGACACCAGAAGTCAACC	GGGTGTAGACGAACCGGATG
ITGA5	CAACTTCTCCTTGGACCCCC	GTCCTCTATCCGGCTCTTGC
Fibronectin	GTATACGAGGGCCAGCTCAT	CCCAGGAGACCACAAAGCTA
NOS2	CGCAGAGAACTCAGCCTCAT	TGCCTTGAGAACTTCGGGAC
TNF α	CTTCTGCCTGCTGCACTTG	GTCACTCGGGGTTGAGAAG
Periostin	ACAAGAAGAGGTCACCAAGGTC	CTTGCAACTTCCTCACGGGT
α -SMA	CCCCATCTATGAGGGCTATG	CAGTGCCATCTCATTTTCA



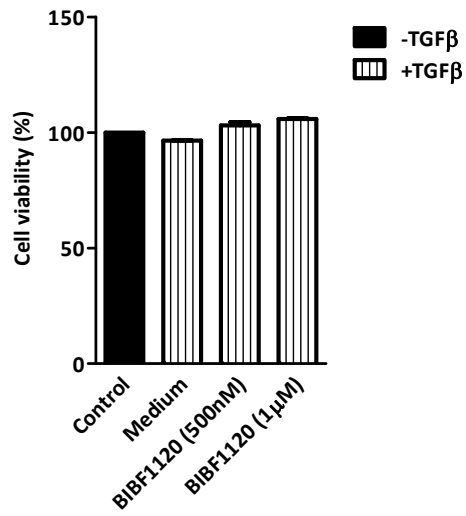
Supplementary Figure S1: Graph depicts % cell viability (as assessed by Alamar blue assay) in 3T3 fibroblasts treated with or without TGFβ (5ng/ml) ± 500nM or 1µM tyrosine kinase inhibitor BIBF1120. Bars represent mean ± SEM, n=3.



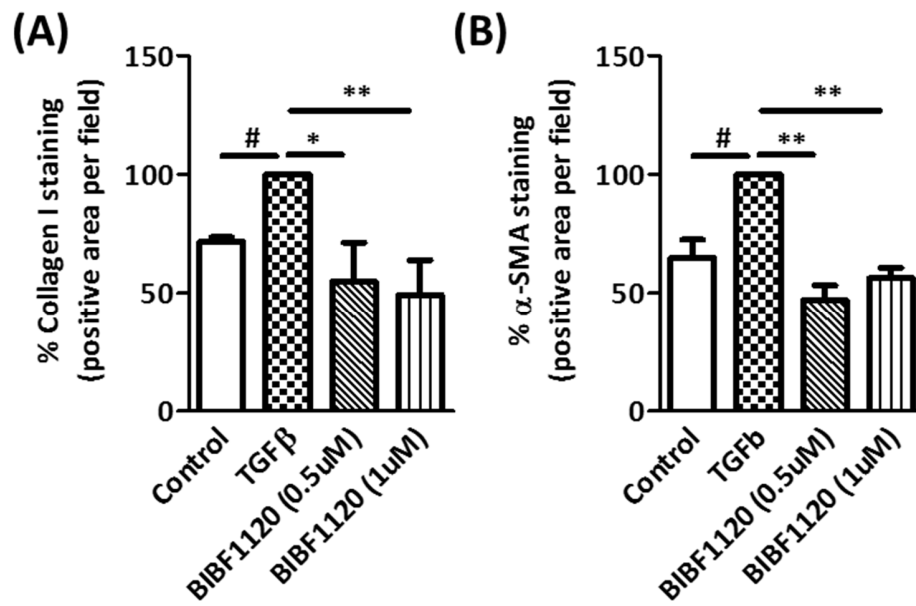
Supplementary Figure S2: Graph depicts quantitative immunostaining analysis of α -SMA (**A**) and Collagen I (**B**) stained LX2 cells that were treated with or without TGF β (5ng/ml) \pm 500nM or 1 μ M tyrosine kinase inhibitor BIBF1120. Bars represent mean \pm SEM, n=3. #p<0.05 denotes significance versus control LX2 cells. *p<0.05, **p<0.01 denotes significance versus TGF β -treated LX2 cells.



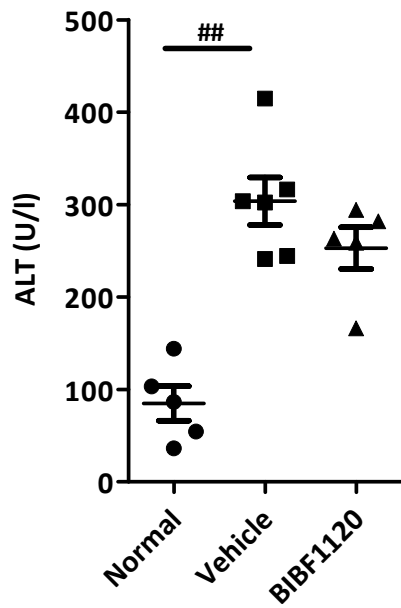
Supplementary Figure S3: **(A)** Respective western blot and **(B, C)** quantitative band intensity analysis (normalized with respective β -actin bands and expressed in %) depicting Collagen-I, ITGA5, pFAK and β -actin protein expression in human HSCs (LX2 cells) treated with or without TGF β (5ng/ml) \pm 1 μ M tyrosine kinase inhibitor BIBF1120. Bars represent mean \pm SEM, n=3. ##p<0.01 denotes significance versus control cells; **p<0.01 denotes significance versus TGF β -treated cells.



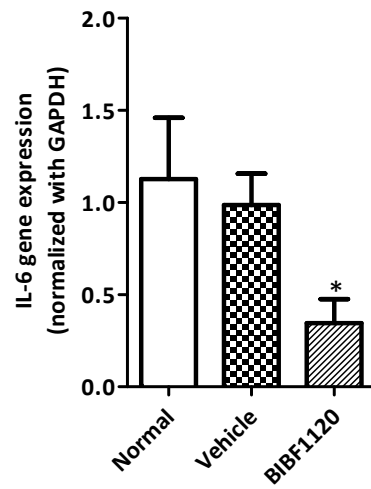
Supplementary Figure S4: Graph depicts % cell viability (as assessed by Alamar blue assay) in human LX2 cells treated with or without TGFβ (5ng/ml) ± 500nM or 1μM tyrosine kinase inhibitor BIBF1120. Bars represent mean ± SEM, n=3.



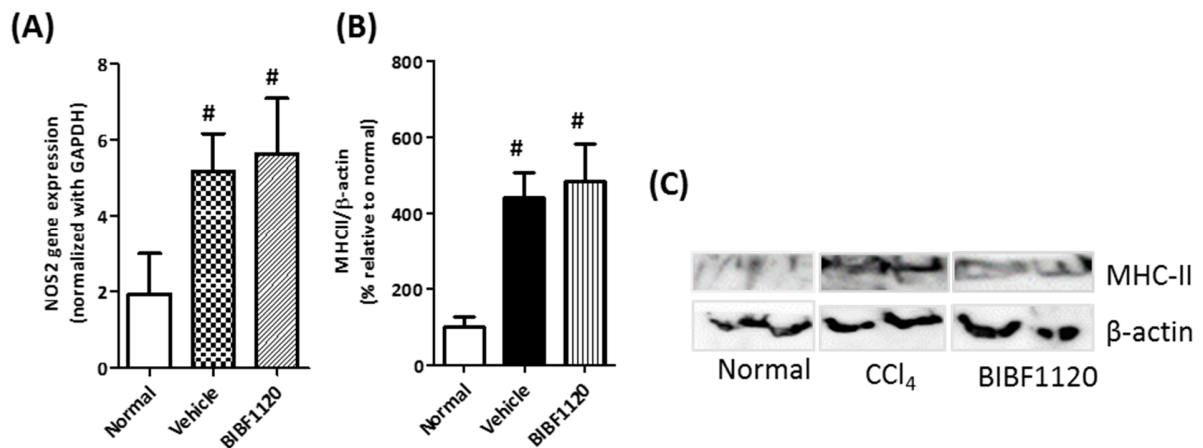
Supplementary Figure S5: Graph depicts quantitative immunostaining analysis of Collagen I and α -SMA stained primary human hepatic stellate cells (pHSCs) that were treated with or without TGF β (5ng/ml) \pm 500nM or 1 μ M tyrosine kinase inhibitor BIBF1120. Bars represent mean \pm SEM, n=3. #p<0.05 denotes significance versus control pHSCs. *p<0.05, **p<0.01 denotes significance versus TGF β -treated pHSCs.



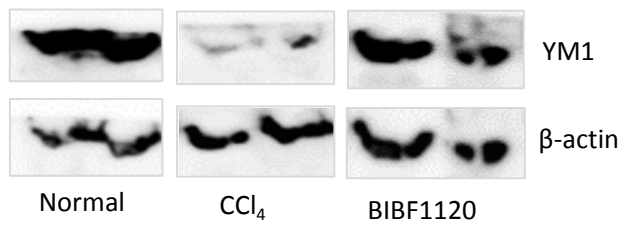
Supplementary Figure S6: Alanine aminotransferases (ALT) levels (expressed as U/l) as analyzed in the serum from olive-oil-treated controls (normal), vehicle-treated CCl₄ mice (Vehicle) and BIBF1120-treated CCl₄ mice. Each symbol represents individual mice. Bars represent mean ± SEM of n=5 mice per group. ##p<0.01 denotes significance versus olive-oil treated control group.



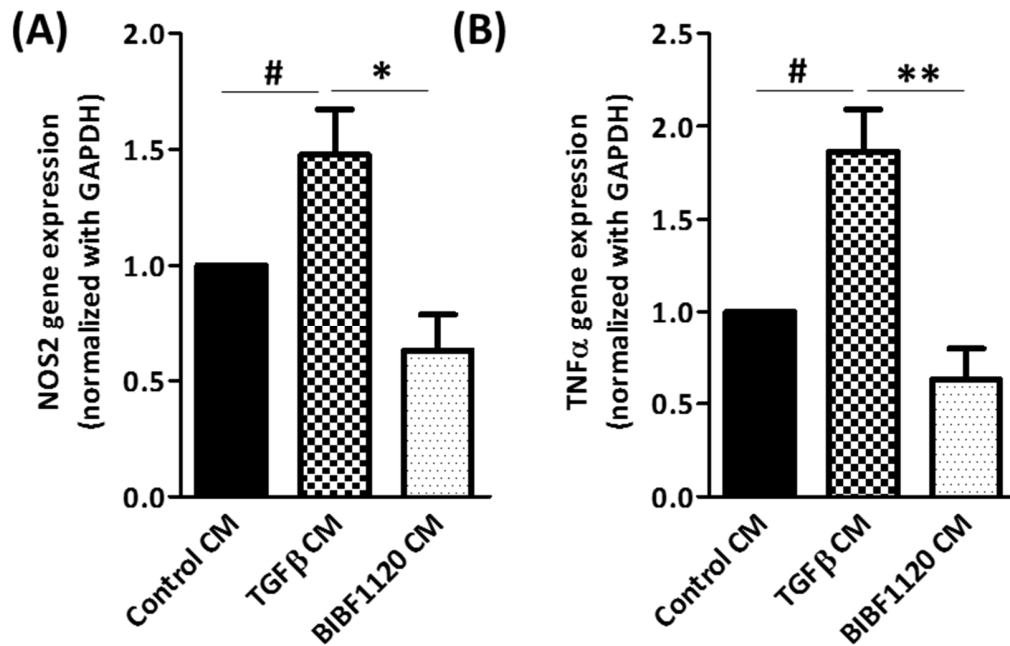
Supplementary Figure S7: Quantitative gene expression (normalized with GAPDH) of IL-6 in the livers of olive-oil-treated controls (normal), vehicle-treated CCl₄ mice and BIBF1120-treated CCl₄ mice. Bars represent mean \pm SEM of n=5 mice per group. *p<0.05 denotes significance versus CCl₄-treated vehicle group.



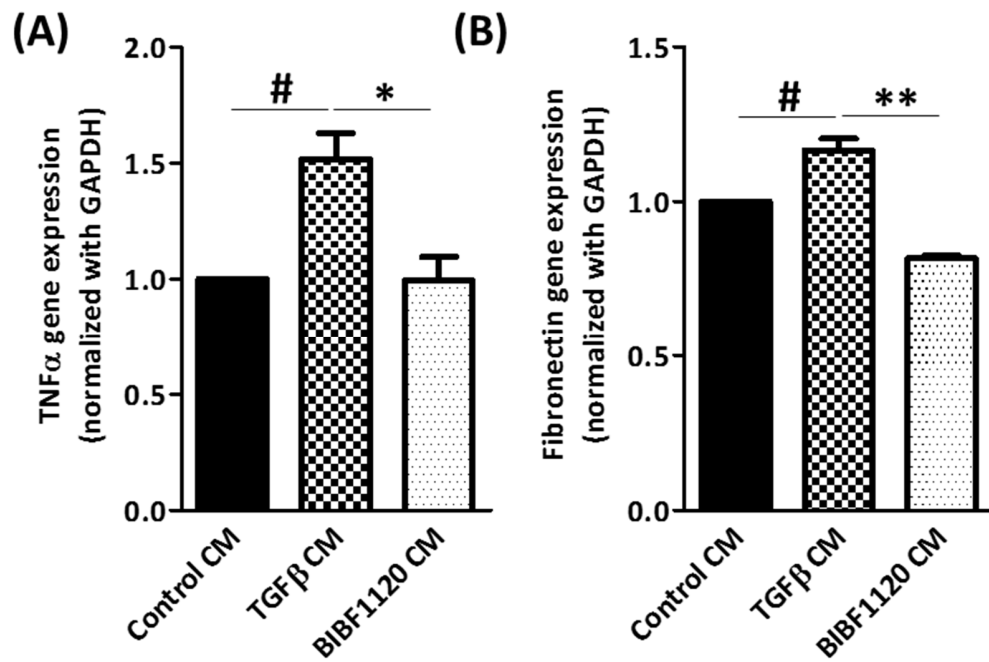
Supplementary Figure S8: (A) Quantitative gene expression (normalized with GAPDH) of NOS2 or iNOS in the livers of olive-oil-treated controls (normal), vehicle-treated CCl₄ mice and BIBF1120-treated CCl₄ mice. **(B)** Quantitative band intensity analysis (normalized with respective β-actin bands and expressed in %) and **(C)** Respective western blot depicting MHC-II and β-actin protein expression in olive-oil-treated controls (normal), vehicle-treated CCl₄ mice and BIBF1120-treated CCl₄ mice. Bars represent mean ± SEM of n=5 mice per group. #p<0.05, ##p<0.01 denotes significance versus olive-oil treated control group.



Supplementary Figure S9: Respective western blot depicting YM1 and β -actin protein expression in olive-oil-treated controls (normal), vehicle-treated CCl₄ mice and BIBF1120-treated CCl₄ mice.



Supplementary Figure S10. Quantitative gene expression (normalized with GAPDH) of M1 macrophage marker, iNOS or NOS2 (nitric oxide synthase 2) **(A)** and inflammation marker TNFα **(B)** analyzed in primary human kupffer cells incubated with conditioned medium from primary human hepatic stellate cells that are treated with medium alone (control CM), TGFβ (5ng/ml) with or without 1μM BIBF1120. Bars represent mean ± SEM of n=3 independent experiments. #p<0.05 denotes significance versus control CM; *p<0.05, **p<0.01 denotes significance versus TGFβ CM.



Supplementary Figure S11. Quantitative gene expression (normalized with GAPDH) of inflammation marker TNF α (A) and fibronectin (B) analyzed in primary human LSECs incubated with conditioned medium from primary human hepatic stellate cells that are treated with medium alone (control CM), TGF β (5ng/ml) with or without 1 μ M BIBF1120. Bars represent mean \pm SEM of n=3 independent experiments. #p<0.05 denotes significance versus control CM; *p<0.05, **p<0.01 denotes significance versus TGF β CM.