Correlations of factors

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1 Computations

1.1 Correlations

Four different sets of correlations are computed, with different advantages and disadvantages. First, correlation of time point averages are computed. These correlations reflect the general time development of the factors. If a correlation coefficient is high, the factors share a similar time profile. The correlations all reflect vectors of 8 values, and are comparable among each other. This correlation is denoted with Av.

Second, correlations for indivdual mice are computed. The time point is ignored here. These correlations reflect a possible mechanistic relation of the factors. The correlations reflect long vectors of variable length because not all factors are measured in each mouse. Correlations which include one of SMA, CTFG, S1004A cell count are at about 24 values, the others at about 40 values. Thus, the correlation values are not completely comparable, especially those including the SMA, CTFG, S1004A count. This correlation is denoted with All. The disadvantage is that it completely ignores the time from duct ligation.

2 1 COMPUTATIONS

Third, correlations for individual mice by each individual time point are computed. These correlations reflect a possible mechanistic relation best. However, the correlations are computed for short vectors of up to 5 values. For some correlations including the SMA, CTFG, S1004A count, there are only 2 values (thus, the correlation can only 1 or -1 in this case). There are eight correlations of this type and they are deonted with T0h, ..., T14d. The main disadvantage is the low robustness due to the small number of values.

Fourth, correlations for indivdual mice for two or three adjacent time points are computed. The correlations are computed for vectors from 6 to 15 values and are more robust than the single time point correlation above. They also represent time developments in a restricted time range. There are eleven correlations of this type. The time frames which coincide with the disease phases are denoted with Init (6-12h), Perp (18f-2d), and Progr (5-14d). The other correlations are simply denoted by the time frame: 6-18h, 12-18h, 12-30h, 18-30h, 30h-2d, 30h-5d, 2-5d, and 2-14d. The correlation show the correlation at a specific time frame and the main disadvantage is that there are quite many correlation numbers to consider. This problem is tackled with the computation of a consensus score.

1.2 Consensus correlations

The following observations are based on the consensus of correlation. The consensus is calculated from the correlations shown above, where the correlation among all mice, and the correlation of time averages has 4 times higher weight:

$$C(\alpha, \beta) = \max \left\{ \sum_{i=1}^{21} w_i s_i \operatorname{pos} \left(\operatorname{corr}_i(\alpha, \beta) \right), \sum_{i=1}^{21} w_i s_i \operatorname{neg} \left(\operatorname{corr}_i(\alpha, \beta) \right) \right\}$$

$$w_{1,2} = 4, w_{i>3} = 1$$

$$\operatorname{corr}_i(\alpha, \beta) \dots \operatorname{PEARSON} \text{ correlation } \# i \text{ of factors } \alpha \text{ and } \beta$$

$$\operatorname{pos}(x) = \begin{cases} x, & \text{if } x > 0 \\ 0, & \text{if } x \leq 0 \end{cases} \dots \operatorname{positive part}$$

$$\operatorname{neg}(x) = \begin{cases} 0 & \text{if } x > 0 \\ -x & \text{if } x \leq 0 \end{cases} \dots \operatorname{positive part}$$

$$s_i = \min \left\{ 1, -\log_{100} p_i(\alpha, \beta) \right\}$$

$$p_i(\alpha, \beta) \dots \text{t-test significance of } \operatorname{corr}_i(\alpha, \beta)! = 0$$

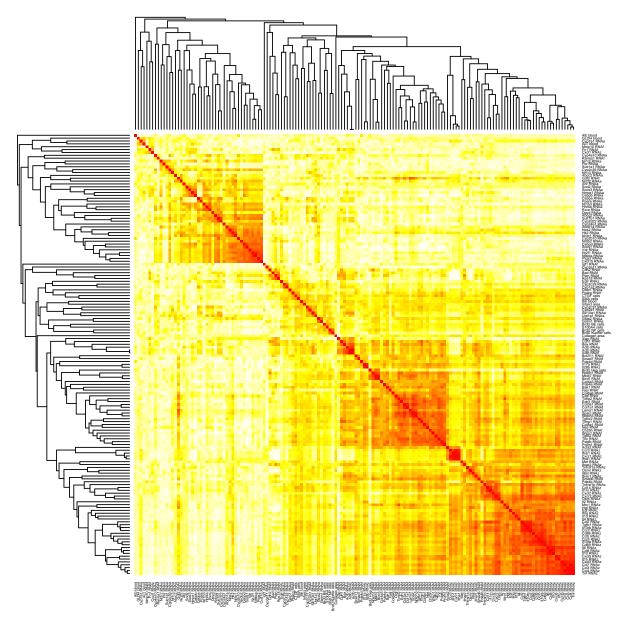
The square of the correlation is chosen to emphasize the differences among relatively high (>0.5) correlations. rank_i refers to the list of the correlations of all factors ordered by their absolute values. The exponents is chosen such that the rank modification has a smaller contribution than the correlation itself. The rank modification help to balance factors with an uneven distribution of correlations.

The following tables show factors with the highest consensus correlation (and anti-correlation). They are the main candidates for marker genes.

The following abbreviations are used:

RNAa	RNA of the ADME set
RNAz	TNA of the zytokine set
RNAf	RNA of the fibrosis set
Init	correlation of mice of the initial phase (6-12h)
Perp	correlation of mice of the initial phase (18h-2d)
Progr	correlation of mice of the initial phase (5-14d)
All	correlation of all individual mice
Av	correlation of the time averages
T0h14d	correlation of the mice of a single time point

2 Heatmaps of consensus correlations



 ${\bf Figure} \ {\bf 1} - {\bf Heatmap} \ {\bf of} \ {\bf consensus} \ {\bf correlation} \ {\bf of} \ {\bf all} \ {\bf factors}.$

$2.1 \quad Heatmap \ of \ 80 \ cluster \ set$

 ${\bf Table}~{\bf 1}-{\rm cluster}~{\rm definition}~{\rm of}~80~{\rm cluster}~{\rm set}$

Cluster name	factors
Actb RNAf	Actb RNAf, Actb RNAz, Actb RNAa, Bax RNAf
Cytokines	CcI4 RNAz, Tnf RNAz, CcI7 RNAz, CcI3 RNAz, II10 RNAz, CcI8 RNAz, II6 RNAz, Cd69 RNAz, II10ra RNAz, Ccr5 RNAz, CcI5
	RNAz, II10rb RNAz, Ccr2 RNAz, Cxcl3 RNAz, Tgfb1 RNAz, Cd86 RNAz, II1b RNAz, Cxcl5 RNAz, Ccl2 RNAz, II13 RNAz, Ifng
	RNAz, II4 RNAz, Egf RNAz
Cxcr1 RNAz	Cxcr1 RNAz, Ifna1 RNAz, Ccr3 RNAz, Ifnb1 RNAz
Birc5 RNAf	Birc5 RNAf, Mki67 RNAf, Notch1 RNAf
Fibrosis genes	Sparc RNAf, Notch3 RNAf, Col1a1 RNAf, Col3a1 RNAf, Lama1 RNAf, Pdgfb RNAf, Prom1 RNAf, Timp2 RNAf, Edn1 RNAf, Tgfb2
ŭ	RNAf, Tnc RNAf, Col8a1 RNAf, Wisp1 RNAf, Tgfbr2 RNAf, Timp1 RNAf, Ctgf RNAf, Nes RNAf, Ch25h RNAf
II28b RNAz	II28b RNAz, II17a RNAz
ADME genes	Nfkbia RNAa, Nr2f1 RNAa, Cxcl15 RNAa, Ptgs2 RNAa, Cebpd RNAa, Socs1 RNAa, Vdr RNAa, Nr0b2 RNAa
II1rn RŇAz	Il1rn RNAz, Cd14 RNAz, Osm RNAz, Cxcr2 RNAz, Cxcl2 RNAz, Tnfrsf1b RNAz
Sult1b1 RNAa	Sult1b1 RNAa, Cyp2d22 RNAa
Nr3c1 RNAa	Nr3c1 RNAa, Hk2 RNAa, Nos2 RNAa
Ifnar1 RNAz	Ifnar1 RNAz
Hgf RNAz	Hgf RNAz, Mrc1 RNAz
Gdf2 RNAf	Gdf2 RNAf
Osmr RNAz	Osmr RNAz, Il6st RNAz, Tnfrsf1a RNAz
Bak1 RNAf	Bak1 RNAf
Cdh1 RNAf	Cdh1 RNAf
Cyp1a2 RNAa	Cyp1a2 RNAa, Cyp2c29 RNAa
Rxra RNAa	Rxra RNAa, Hnf4a RNAa, Nr1h3 RNAa
Pten RNAf	Pten RNAf, Rps18 RNAf
Pde4b RNAf	Pde4b RNAf
Hmox1 RNAa	Hmox1 RNAa, Cebpb RNAa
Cyp2b10 RNAa	Cyp2b10 RNAa
II2 RNAz	II2 RNAz
BrdU Kup cells	BrdU Kup cells
Gstm1 RNAa	Gstm1 RNAa, Gsta2 RNAa
Bcl2l11 RNAf	Bcl2l11 RNAf
BrdU Hep cells	BrdU Hep cells
Abcg2 RNAa	Abcg2 RNAa
Pde4a RNAf	Pde4a RNAf
Smad6 RNAf	Smad6 RNAf
Nr2f2 RNAa	Nr2f2 RNAa
Ppara RNAa	Ppara RNAa, Cebpa RNAa
Col6a6 RNAf	Col6a6 RNAf
Socs3 RNAa	Socs3 RNAa
Fasl RNAf	Fasl RNAf
Abcb1a RNAa	Abcb1a RNAa
Acta2 RNAf	Acta2 RNAf
Dpyd RNAa	Dpyd RNAa
Cdh2 RNAf	Cdh2 RNAf
Ifnar2 RNAz	Ifnar2 RNAz
Col4a3 RNAf	Col4a3 RNAf
Cyp2c37 RNAa	Cyp2c37 RNAa
Cyp2e1 RNAf	Cyp2e1 RNAf
Cyp24a1 RNAa	
Slc10a1 RNAa	Sic10a1 RNAa
Ugt1a1 RNAa	Ugt1a1 RNAa
Bad RNAf	Bad RNAf
Met RNAz Smad7 RNAf	Met RNAz Smad7 RNAf
SMA cells	SMA cells
CTGF cells	CTGF cells
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cluster definition of 80 cluster set- continued

Cluster name	factors
Sod2 RNAa	Sod2 RNAa
Xiap RNAf	Xiap RNAf
Cyp2c39 RNAa	Cyp2c39 RNAa
Ahr RNAa	Ahr RNAa
Nr1i3 RNAa	Nr1i3 RNAa
Nr1i2 RNAa	Nr1i2 RNAa
Pparg RNAf	Pparg RNAf
BrdU BE cells	BrdU BE cells
S100A4 cells	S100A4 cells
Abcc2 RNAa	Abcc2 RNAa
Bili blood	Bili blood
Infarct area	Infarct area
Pde4d RNAf	Pde4d RNAf
ALT blood	ALT blood
Cyp7a1 RNAa	Cyp7a1 RNAa
Sult1a1 RNAa	Sult1a1 RNAa
• •	Cyp4a10 RNAa
Por RNAa	Por RNAa
Fn1 RNAf	Fn1 RNAf
Cxcl1 RNAz	Cxcl1 RNAz
Egfr RNAz	Egfr RNAz
Gstp1 RNAa	Gstp1 RNAa
GLDH blood	GLDH blood
Collagen area	Collagen area
Mmp10 RNAf	Mmp10 RNAf
Rarres1 RNAf	Rarres1 RNAf
lgf1 RNAf	Igf1 RNAf
Alb blood	Alb blood
Cyp3a11 RNAa	Cyp3a11 RNAa

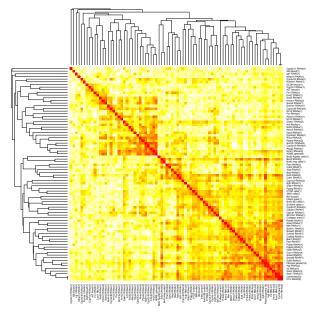


Figure 2 – Heatmap of consensus correlation of 80 clusters set. The colors are scaled to the highest non-diagonal value.

2.2 Heatmap of 30 cluster set

 ${\bf Table} \ {\bf 2} - {\bf cluster} \ {\bf definition} \ {\bf of} \ {\bf 30} \ {\bf cluster} \ {\bf set}$

Cluster name	factors
II28b RNAz	II28b RNAz, II17a RNAz, Pde4d RNAf
ADME	Nfkbia RNAa, Cxcl15 RNAa, Vdr RNAa, Nr2f1 RNAa, Ptgs2 RNAa, Sult1b1 RNAa, Cyp2d22 RNAa, Nr0b2 RNAa, Nr3c1 RNAa,
	Hk2 RNAa, Cebpd RNAa, Socs1 RNAa, Nos2 RNAa, Cyp2b10 RNAa, Cyp24a1 RNAa, Abcg2 RNAa, Rxra RNAa, Hnf4a RNAa,
	Abcb1a RNAa, Nr1h3 RNAa, Dpyd RNAa, Ahr RNAa
Cytokine	Ifng RNAz, II4 RNAz, II10 RNAz, Ccl8 RNAz, Tnf RNAz, II6 RNAz, II10ra RNAz, Ccr5 RNAz, Ccl4 RNAz, Ifnb1 RNAz, Cxcr1 RNAz,
	Ccr3 RNAz, Egf RNAz, Osm RNAz, Cd69 RNAz, Tgfb1 RNAz, Il10rb RNAz, Ccl3 RNAz, Ccl7 RNAz, Il13 RNAz, Ifna1 RNAz, Ccl5
	RNAz, Ccr2 RNAz, Ccl2 RNAz, Il2 RNAz, Il1b RNAz, Cxcl3 RNAz, Cd86 RNAz, Cxcr2 RNAz, Il1rn RNAz, Cd14 RNAz, Cxcl5 RNAz,
	Tnfrsf1b RNAz, Osmr RNAz, Hgf RNAz, Mrc1 RNAz, Cxcl2 RNAz, Pde4b RNAf, Met RNAz
Fibrosis	Birc5 RNAf, Mki67 RNAf, Sparc RNAf, Col3a1 RNAf, Notch3 RNAf, Col1a1 RNAf, Pdgfb RNAf, Tnc RNAf, Col8a1 RNAf, Prom1
	RNAf, Wisp1 RNAf, Timp1 RNAf, Timp2 RNAf, Tgfbr2 RNAf, Lama1 RNAf, Tgfb2 RNAf, CTGF cells, Edn1 RNAf, Ifnar1 RNAz, Ctgf
	RNAf, SMA cells, Nes RNAf, Notch1 RNAf, Ch25h RNAf, Cdh1 RNAf, Bak1 RNAf, BrdU Hep cells, Smad6 RNAf, Col6a6 RNAf,
	Pde4a RNAf, Fasl RNAf, Col4a3 RNAf, Ifnar2 RNAz, Acta2 RNAf
Actb RNAf	Actb RNAf, Actb RNAz, Actb RNAa, Bax RNAf, Xiap RNAf
BrdU Kup cells	BrdU NP cells, BrdU Kup cells, BrdU BE cells, S100A4 cells, Gstm1 RNAa, Gsta2 RNAa
Hmox1 RNAa	Hmox1 RNAa, Cebpb RNAa, Socs3 RNAa, Sod2 RNAa, Cxcl1 RNAz
Gdf2 RNAf	Gdf2 RNAf
Cyp1a2 RNAa	Cyp1a2 RNAa, Cyp2c29 RNAa
Tnfrsf1a RNAz	Tnfrsf1a RNAz, II6st RNAz
Cyp2c39 RNAa Fn1 RNAf	Cyp2c39 RNAa, Nr1i3 RNAa Pten RNAf, Rps18 RNAf, Bad RNAf, Cdh2 RNAf, Fn1 RNAf
Pparg RNAf	Pparg RNAf
Cyp2c37 RNAa	· •
Nr2f2 RNAa	Nr2f2 RNAa, Abcc2 RNAa
Ppara RNAa	Ppara RNAa, Cebpa RNAa
Bcl2l11 RNAf	Bcl2l11 RNAf, Smad7 RNAf
Egfr RNAz	Egfr RNAz
Slc10a1 RNAa	Slc10a1 RNAa, Ugt1a1 RNAa, Por RNAa
Rarres1 RNAf	Rarres1 RNAf
Gstp1 RNAa	Gstp1 RNAa
Bili blood	Bili blood, Infarct area
GLDH blood	GLDH blood
Nr1i2 RNAa	Nr1i2 RNAa, Sult1a1 RNAa
ALT blood	Cyp7a1 RNAa, ALT blood
• •	Cyp3a11 RNAa
Collagen area	Collagen area
Mmp10 RNAf	Mmp10 RNAf
Alb blood	Alb blood
Igf1 RNAf	Igf1 RNAf

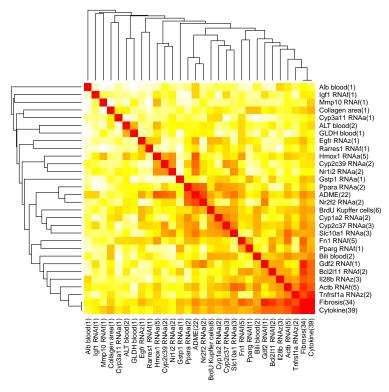


Figure 3 – Heatmap of consensus correlation of 80 clusters set.

2.3 Heatmap of main factors

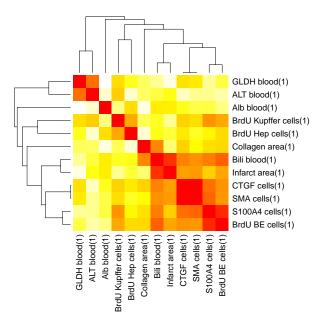
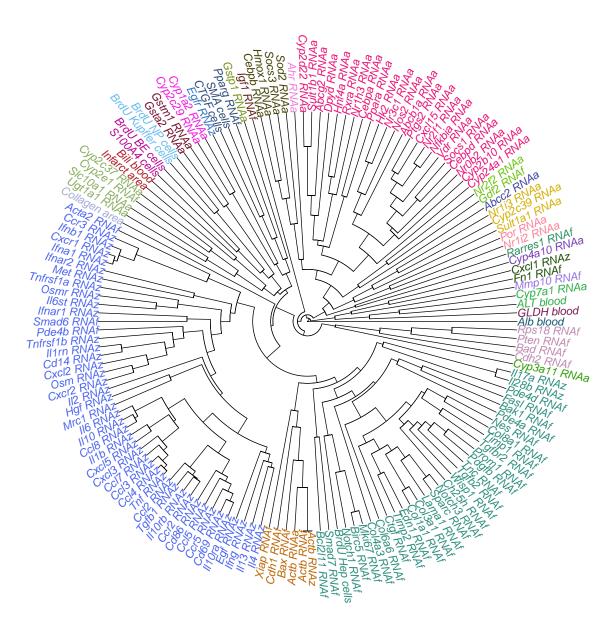
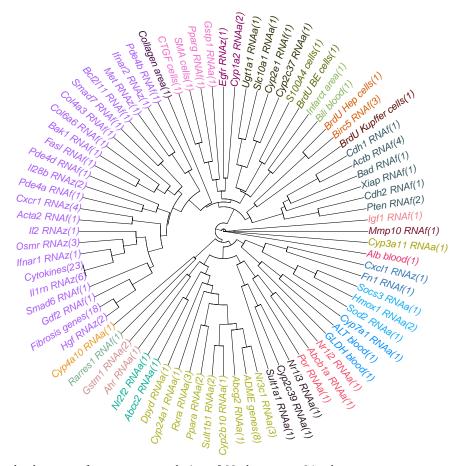


Figure 4 – Heatmap of consensus correlation of main factors.

3 Cluster maps of consensus correlations



 ${\bf Figure~5-Circular~dendrogram~of~consensus~correlation~of~all~factors,~30~colors.}$



 ${\bf Figure~6}-{\rm Circular~dendrogram~of~consensus~correlation~of~80~cluster~set,~24~colors.}$

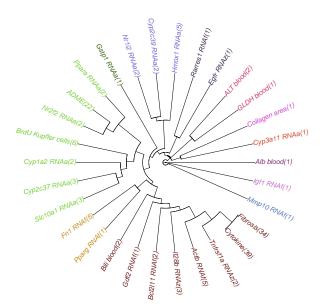


Figure 7 – Circular dendrogram of consensus correlation of 30 cluster set, 12 colors.

4 Most correlated factors for each main factor

The following digrams are larger versions of the respective diagrams in the main manuscript. Here, the list of 18 most correlated factors is shown. Each entry holds the respective number.

4.1 ALT blood

Factor	Cons		All				12-30h											T18h			T5d	T14d
Cyp7a1 RNAa	9.12		-0.66				-0.54									-0.96		-0.36		-0.81	-0.98	
		0.02	0	0.62	0.28	0.94	0.11			0.01			0.005			0.01	0.27		0.06	0.1		0.12
Hmox1 RNAa	8.62		0.62	0.29	0.7	0.47	0.02	0.39	0.87		0.22	0.81	0.8	0.74	-0.73		-0.22		0.6	0.98	0.58	0.5
		0.04	0	0.41	0.02	0.17	0.96		0.001		0.57		0.001	0.02	0.16	0.19	0.72	0.12		0.003	0.42	0.39
GLDH blood	8.46	0.76	0.67	0.47	0.31	0.25	0.9		0.43	0.82	0.67	0.15	0.46	0.42	0.59	0.34	0.83	0.21	0.95	0.89		-0.28
6 1 1 5014	0.75	0.03	0	0.17	0.39		0.0004			0.004		0.68	0.09	0.23	0.29	0.58	0.08	0.74	0.01	0.04	0.68	0.65
Cebpb RNAa	6.75	0.72	0.49	0.22	0.64		-0.11	0.37		0.49	0.04	0.47	0.5	0.39	0.34		-0.67		-0.09		0.14	
C II DNIA	0.75		0.002	0.54	0.05	0.28	0.76	0.3	0.003		0.92	0.2	0.07	0.3	0.57	0.25	0.22	0.06	0.88	0.1	0.86	0.91
Cxcl1 RNAz	6.75		0.5		0.43		-0.13	0.37	0.63	0.33	0.47	0.55	0.52	0.53		-0.58		0.8	0.11	0.49		-0.04
E 4 DNAC	0.40		0.001	0.07	0.21	0.21	0.71	0.29	0.05	0.35	0.2	0.12	0.05	0.15	0.03	0.3	0.32	0.11	0.85	0.4	0.11	0.95
Fn1 RNAf	6.42		0.6	-0.18				0.12	0.61	0.49	0.41	0.59	0.64	0.61		-0.37			0.29	0.7	0.63	0.44
C 0 DNA	F 07	0.02	0	0.61	0.57	0.86	0.68	0.74	0.06	0.15	0.27	0.1	0.01	0.08	0.48	0.54	0.23	0.48	0.64	0.19	0.37	0.46
Socs3 RNAa	5.97	0.51	0.46	0.09	0.31		-0.18	0.19	0.75	0.5	0.25	0.68	0.63	0.37	-0.01		-0.43		-0.2	0.87		-0.36
C 0.100 DNIA	E 0E		0.004	0.81	0.38	0.46	0.62	0.6	0.01	0.15	0.52	0.04	0.02	0.32	0.98	0.46	0.47	0.36	0.75	0.06	0.49	0.55
Cyp2d22 RNAa	5.95	-0.62			0.002			-0.35	-0.6	-0.67			-0.53		-0.64		0.45	0.17		-0.85		-
A .I DNIAC	F 0F	0.1	0.01	0.93	1	0.54	0.47	0.32	0.07	0.03	0.22	0.06	0.05	0.36	0.24	0.7	0.44	0.78	0.29	0.07	0.66	0.43
Actb RNAf	5.95	0.25	0.39	0.1	0.25	0.18	0.1	0.17	0.76	0.48	0.41	0.89	0.67	0.66		-0.14		0.45	0.12	0.91	0.92	0.84
Calara DNA -	E 01	0.55	0.01	0.78	0.49	0.62	0.78	0.65	0.01	0.16		0.001	0.01	0.05	0.57	0.82	0.92	0.44	0.85	0.03	0.08	0.08
Cebpa RNAa	5.91			-0.41		0.42	-0.7				-0.69		-0.31	-0.15		-0.65				-0.52		
If-L1 DNA-	E 06		0.002	0.24	0.49	0.23	0.02	0.13	0.97	0.15	0.04	0.5	0.28	0.7	0.43	0.23	0.8	0.12	0.05	0.37	0.62	0.97
Ifnb1 RNAz	5.60					-0.38			-	-0.42 0.23	1 1			-0.58		-0.22			-0.9		-0.43	
II1 um DNA =	E 70	0.33	0.04	0.42	0.34	0.28	0.02	0.04	0.7		0.4	0.72	0.11	0.1		0.73	0.62	0.48	0.04	0.75	0.57	0.97
II1rn RNAz	5.79	-0.02 0.96	0.28	0.29	0.61	0.64	0.51	0.48	0.8	0.56	0.18	0.79	0.38	-0.01 0.97	-0.69 0.2	0.13	0.53	0.8	0.54	0.93	0.49	0.62
Ppara RNAa	5.74	-0.5	-0.41	-0.4			•		-0.18		-0.4	-0.5	-0.51	-0.39		-0.54					-0.39	
rpara NivAa	5.74	0.21	0.01	0.26	0.37	0.98	0.28	0.2	0.61	0.27	0.29	0.17	0.06	0.29		0.35	0.88	0.13	0.07	0.08	0.61	0.33
Cxcl2 RNAz	5.67		0.3	0.20	0.63	0.68	0.26	0.2	0.74	0.78	0.24	0.17		-0.27		0.33	0.31	0.84	0.89	0.89	0.15	
CXCIZ INIAZ	5.07	0.76	0.06	0.3	0.05	0.03	0.31	0.14	0.74	0.70	0.53	0.30	0.96	0.48	0.64	0.62	0.61	0.04	0.05	0.03	0.15	0.99
Actb RNAa	5.64		0.34	0.36	0.47	0.19	0.05	0.09	0.69	0.43	0.33	0.86	0.62	0.40	-0.48		0.15		-0.03	0.89	0.86	0.86
/ ICLD TYTA	0.04	0.75	0.04	0.3	0.17	0.59	0.88	0.8	0.03	0.40		0.003		0.08		0.31	0.81	0.46	0.96	0.04	0.14	0.06
Slc10a1 RNAa	5 62	-0.32						-0.44	-0.44				-0.57			-0.76					-0.85	
Sicrour raina	0.02	0.44	0.05	0.75	0.16	0.91	0.72	0.2	0.2	0.08	0.64	0.02	0.03	0.04	0.5	0.14	0.57	0.65	0.19	0.02	0.15	0.12
Pde4b RNAf	5.54	-0.15		0.23	0.66	0.64	0.11	0.45	0.74	0.48	0.42	0.82	0.33	0.15		0.52	-0.01	0.85	0.19	0.78	0.10	0.35
. 20.2 100	0.01	0.72	0.12	0.53	0.04	0.05	0.76	0.19	0.01	0.17	0.27	0.01	0.24	0.7	0.37	0.37	0.99	0.07	0.76	0.12	0.1	0.56
Ch25h RNAf	5.51	-0.07		0.15	0.44	0.4	0.27	0.37	0.79	0.61	0.46	0.81	0.33	0.41	0.74	0.23	0.16	0.52	0.29	0.99	0.91	0.57
2	0.01	0.87	0.15	0.67	0.2	0.25	0.45	0.29	0.01	0.06		0.01		0.27	0.15	0.7	0.8	0.37		0.002		0.32

4.2 GLDH blood 11

4.2 GLDH blood

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
ALT blood	8.46	0.76	0.67	0.47	0.31	0.25	0.9	0.72	0.43	0.82	0.67	0.15	0.46	0.42	0.59	0.34	0.83	0.21	0.95	0.89	-0.26	-0.28
		0.03	0	0.17	0.39	0.48	0.0004	0.02	0.22	0.004	0.03	0.68	0.09	0.23	0.29	0.58	0.08	0.74	0.01	0.04	0.68	0.65
Cyp7a1 RNAa	8.01	-0.74	-0.56	0.11	-0.27	-0.29	-0.62	-0.83	-0.44					-0.41	0.23	-0.45	0.56	-0.68	-0.88	-0.73	0.09	0.52
			0.0002	0.77	0.46		0.06						0.01		0.71	0.45					0.91	0.37
CTGF cells	6.81	-0.43		0.21	0.73					-0.48					-0.36		0.69		-0.97		-0.99	-1
		0.28	0.07	0.69	0.1					0.34				0.003		0.55		0.34				
BrdU NP cells	6.77		-0.42	-0.18						-0.5							-0.08				-0.84	
		0.11	0.01	0.61	0.14		0.26	0.05		0.14		0.09	0.29	0.77	0.8	0.74	0.9	0.39	0.1		0.07	
Fn1 RNAf	6.76		0.6	0.03	0.8		-0.18			0.26				0.19	-0.12		-0.54			0.67		-0.47
C144 !!	0.00	0.01	0	0.93	0.01					0.47			0.15			0.62	0.34		0.4	0.21	0.9	0.42
SMA cells	6.63	-0.35	-0.39	0.19	0.37					-0.56						0.51	0.97		-1	0.93		0.5
M 1 DNA	0.54	0.4	0.06	0.72	0.47					0.25		0.8	0.15				0.14					0.66
Mrc1 RNAz	0.51	-0.64								-0.42					-0.52				-0.31	-0.4	0.96	
T: O DNA£	6 10		0.003	0.92		0.27		0.32			0.15		0.01	0.9	0.37	0.9	0.65	0.9	0.62	0.5	0.04	
Timp2 RNAf	0.13	-0.64 0.09	0.01					0.51	0.1	-0.19 0.61		0.09	-0.68 0.01	-0.4 0.28	0.75		0.44	-0.44 0.45	0.01		-0.25 0.75	
Lama1 RNAf	6.03		-0.49		-0.64					-0.05					-0.74			-0.45				-0.53
Laillai INIAI	0.03		0.001		0.05			0.02	0.09	0.03		0.14		0.85	0.15		0.1	0.43				0.36
BrdU Kup cells	5 00									-0.48			0.28		-0.47						-0.06	
Dido Rup cens	0.00		0.04	0.06		0.04		0.11		0.19					0.43		0.02				0.92	
Gdf2 RNAf	5.97	-0.77			-0.72					0.38							0.64				0.28	
Guiz Turi	0.07		0.001	0.46		0.33				0.27					0.6	0.18			0.29	0.9	0.72	
Sparc RNAf	5.93	-0.61		-0.19		-0.33									-0.43				0.4		-0.31	
	0.00	0.11	-	0.6	0.07		0.19	0.86			0.25		0.01		0.47		0.92	0.61	0.5	0.8		0.53
Gsta2 RNAa	5.92	-0.57	-0.42	-0.29	-0.57	-0.15	0.01	-0.56	-0.78	-0.52	-0.62	-0.07	-0.33	-0.05	-0.75	-0.43	0.83	-0.56	-0.8	-0.07	0.27	0.97
		0.14	0.01	0.41	0.08	0.69	0.98	0.09	0.01	0.12	0.07	0.87	0.25	0.89	0.14	0.47	0.08	0.32	0.1	0.91	0.73	0.01
Sult1a1 RNAa	5.83	0.73	0.46	-0.15	0.16	0.28	0.46	0.16	-0.25	0.16	0.51	0.63	0.71	0.38	-0.59	-0.35	0.46	-0.86	0.41	0.37	0.23	0.71
		0.04	0.003	0.69	0.65	0.43	0.18	0.67	0.48	0.66	0.16	0.07	0.005	0.32	0.29	0.57	0.43	0.06	0.49	0.54	0.77	0.18
Cxcl1 RNAz	5.78	0.89	0.53	0.04	0.53	0.56	-0.1	0.31	0.61	0.03	0.2	0.34	0.35	-0.05	0.78	0.26	-0.42	0.53	-0.12	0.33	-0.11	0.27
		0.003	0.001	0.92	0.12	0.09	0.77	0.39	0.06	0.93	0.61	0.37	0.22	0.9	0.12	0.67	0.49	0.35	0.85	0.59	0.89	0.66
Cebpa RNAa	5.77	-0.45	-0.4	-0.41	-0.75	-0.3	-0.59	-0.67	0.19	-0.29	-0.47	-0.3	-0.29	0.47	-0.86	-0.77	0.54	-0.21	-0.75	-0.13	0.54	0.73
		0.27	0.01	0.24	0.01	0.39	0.07	0.03	0.59	0.42	0.2	0.43	0.31	0.2	0.06	0.12	0.35	0.74	0.14	0.83	0.46	0.16
Notch3 RNAf	5.76	-0.63	-0.41	-0.25	-0.51	-0.4	0.31	-0.04	-0.48	0.31	-0.36	-0.57	-0.67	-0.38	-0.36	-0.11	-0.61	-0.41	0.25	0.46	-0.35	-0.48
		0.09	0.01	0.49	0.13	0.26	0.39	0.9		0.39			0.01			0.86	0.27				0.65	
Birc5 RNAf	5.68	-0.56	-0.35	0.04		-0.65				-0.33			-0.37	-0.08	0.32	0.07	0.03	-0.56	-0.17	-0.16	-0.38	-0.7
		0.15	0.03	0.91	0.21	0.04	0.77	0.37	0.05	0.36	0.21	0.11	0.2	0.84	0.6	0.91	0.96	0.33	0.78	8.0	0.62	0.19

4.3 Alb blood

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B0h-50	d 2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
Ifnb1 RNAz	6.09	0.59	0.27	0.27	0.44	0.66	0.5	0.7	0.6	0.43	0.12	0.02	0.25	0.37	0.31	0.3	-0.21	0.91	0.67	0.33	0.16	0.44
		0.12	0.1	0.44	0.2	0.04	0.14	0.02	0.07	0.21	0.75	0.96	0.39	0.33	0.61	0.63	0.74	0.03	0.22	0.59	0.84	0.45
Ifna1 RNAz	5.57	0.59	0.27	0.46	0.55	0.61	0.42	0.55	0.33	0.25	0.14	0.02	0.22	0.31	0.22	0.54	-0.61	0.65	0.55	-0.03	0.31	0.34
		0.12	0.1	0.18	0.1	0.06	0.23	0.1	0.35	0.49	0.73	0.96	0.46	0.42	0.73	0.35	0.27	0.23	0.34	0.96	0.69	0.58
CTGF cells	5.43	0.76	0.25	-0.23	0.3	0.45	-0.84	-0.34	-0.72	0.75	0.84	0.63	0.43	-0.03	-0.68	0.17	-0.99	-0.87	-0.4	0.63	-0.12	-0.93
		0.03	0.25	0.66	0.56	0.37	0.04	0.51	0.1	0.09	0.04	0.18	0.25	0.95	0.52	0.89	0.1	0.32	0.74	0.57	0.93	0.25
Sult1a1 RNAa	5.25	0.43	0.35	0.15	0.29	0.47	0.47	0.44	0.42	0.57	0.26	0.39	0.17	0.06	-0.11	0.17	0.45	0.03	0.53	0.6	0.88	0.07
		0.29	0.03	0.69	0.41	0.17	0.17	0.2	0.22	0.09	0.49	0.3	0.57	0.87	0.86	0.78	0.45	0.96	0.36	0.28	0.12	0.91
Cyp2d22 RNAa	5.03	-0.31	0.2	0.2	-0.09	0.32	0.44	0.38	0.57	0.7	0.44	0.66	0.56	0.36	0.3	0.26	-0.72	0.11	0.64	0.81	0.62	0.35
		0.45	0.23	0.59	0.79	0.37	0.21	0.27	0.09	0.02	0.23	0.05	0.04	0.34	0.63	0.68	0.17	0.85	0.25	0.09	0.38	0.56
Cyp2c39 RNAa	4.89	0.34	0.36	0.7	0.49	0.38	0.12	0.01	0.42	0.34	-0.01	0.51	0.34	0.17	-0.42	0.74	0.03	-0.13	0.1	0.84	0.69	0.14
		0.41	0.03	0.02	0.15	0.27	0.73	0.98	0.23	0.33	0.99	0.16	0.23	0.65	0.49	0.15	0.96	0.84	0.88	0.08	0.31	0.83
Cyp2b10 RNAa	4.87	-0.19	0.27	0.76	0.31	-0.07	0.01	0.08	0.76	0.6	0.001	0.91	0.7	0.2	-0.49	0.82	0.99	0.45	-0.31	1	0.58	0.29
		0.66	0.1	0.01	0.38	0.85	0.97	0.82	0.01	0.07	1	0.001	0.01	0.6	0.4	0.09	0.0005	0.45	0.61	0	0.42	0.64
Cxcr1 RNAz	4.86	0.61	0.25	0.21	0.41	0.65	0.23	0.47	0.42	0.14	0.11	0.001	0.21	0.3	0.34	0.22	-0.27	0.76	0.31	-0.01	0.05	0.29
		0.11	0.13	0.57	0.24	0.04	0.52	0.17	0.23	0.71	0.78	1	0.47	0.43	0.57	0.72	0.66	0.14	0.62	0.98	0.95	0.64
Cyp1a2 RNAa	4.74	-0.77	-0.35	-0.09	-0.36	-0.47	-0.13	0.23	-0.26	-0.27	-0.06	-0.48	-0.41	0.09	0.67	-0.45	-0.78	0.07	-0.05	-0.63	0.33	0.19
		0.03	0.03	0.81	0.31	0.18	0.72	0.53	0.47	0.45	0.88	0.19	0.14	0.82	0.21	0.45	0.12	0.91	0.94	0.26	0.67	0.76
SMA cells	4.52	0.75	0.21	-0.39	0.41	0.77	-0.77	-0.43	-0.44	-0.24	0.36	0.19	0.52	0.35	0.68	-0.99	-0.75	0.65	-0.68	-0.82	-0.51	0.75
		0.03	0.34	0.44	0.42	0.07	0.08	0.39	0.38	0.64	0.48	0.72	0.15	0.5	0.52	0.1	0.46	0.55	0.53	0.38	0.66	0.46
S100A4 cells	4.26	0.73	0.14	-0.48	0.32	0.79	-0.74	-0.48	-0.62	0.54	0.86	-0.39	-0.2	-0.12	-0.84	-0.64	-0.11	0.84	-0.34	-0.94	0.39	-0.81
		0.04	0.51	0.33	0.54	0.06	0.09	0.33	0.19	0.27	0.03	0.45	0.61	0.82	0.37	0.56	0.93	0.37	0.78	0.21	0.75	0.4
Pde4d RNAf	4.24	0.86	0.31	0.82	0.34	-0.1	0.06	-0.16	-0.3	-0.15	0.09	-0.08	0.18	0.41	0.17	0.84	0.82	-0.47	-0.03	-0.24	0.41	0.64
		0.01	0.06	0.004	0.33	0.77	0.87	0.66	0.41	0.68	0.81	0.83	0.53	0.27	0.79	0.08	0.09	0.43	0.96	0.69	0.59	0.25
Col6a6 RNAf	4.11	0.65	0.05	-0.22	-0.42	-0.55	-0.58	-0.69	-0.67	-0.68	-0.14	-0.16	0.09	0.31	-0.66	-0.31	0.28	-0.65	-0.67	-0.68	0.6	0.44
		0.08	0.75	0.55	0.23	0.1	0.08	0.03	0.03	0.03	0.73	0.69	0.76	0.42	0.22	0.62	0.65	0.24	0.21	0.2	0.4	0.46
Bili blood	4.11	0.77	0.33	0.36	0.17	0.12	0.37	0.04	-0.39	0.15	0.37	0.16	0.14	0.03	0.42	0.92	0.92	-0.58	0.45	-0.3	0.2	-0.1
		0.03	0.04	0.3	0.64	0.75	0.29		0.27		0.29	0.66	0.63	0.93	0.49	0.03	0.03	0.31	0.45	0.63	0.75	0.88
Actb RNAa	4.06	0.94	0.2	0.57	0.46	0.5	0.05	-0.02	-0.59	-0.42	0.07	-0.41	-0.21	0.15	0.88	0.63	0.96	-0.08			0.4	0.25
		0.001	0.22	0.09	0.18	0.14	0.89	0.95	0.07	0.22	0.86	0.27	0.47	0.71	0.05	0.26	0.01	0.9	0.78	0.15	0.6	0.68
Cebpa RNAa	3.97	-0.69	-0.02	0.5	-0.05	-0.83	0.55		0.3	0.54	0.53	0.42	0.48	0.38	0.67	0.77	-0.87	-0.94	0.67	0.78	0.78	0.55
		0.06	0.88	0.14		0.003		0.47	0.41	0.1		0.26	0.08	0.31	0.22					0.12		0.34
Ccr3 RNAz	3.95	0.61	0.24	0.19	0.44	0.58	-0.002	0.36	0.15	-0.1	0.1	-0.03	0.18	0.25	0.38	0.26	-0.77	0.69	0.07	-0.32	-0.17	0.16
		0.11	0.15	0.6	0.2	0.08	0.99	0.31	0.68	0.78	0.81	0.94	0.53	0.51	0.52	0.67	0.12	0.2	0.91	0.6	0.83	0.79
Infarct area	3.9	0.79	0.33	0.14	0.01	-0.01	0.5	0.1	-0.39	0.11	0.31	0.05	0.18	0.28	0	0.96	0.71	-0.49	0.6	-0.3	0.12	0.56
		0.02	0.04	0.7	0.98	0.97	0.14	0.78	0.27	0.75	0.39	0.89	0.51	0.44	n.a.	0.01	0.18	0.4	0.28	0.62	0.85	0.33

4.4 Bili blood

Factor Cons A	v All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
Tnfrsf1b RNAz 10.3 0.	0.85	0.44	0.83	0.81	0.63	0.61	0.58	0.4	0.8	0.79	0.7	0.11	0.05	0.03	0.09	0.88	0.31	0.5	0.44	-0.96
0.0	001 0	0.2	0.003	0.005	0.05	0.06	0.08	0.26	0.01	0.01	0.005	0.77	0.93	0.96	0.89	0.05	0.61	0.4	0.56	0.01
Ccl2 RNAz 10.3 0.		0.45	0.88	0.65	0.62	0.7	0.7	0.61	0.72	0.7	0.64	-0.04		-0.59	-0.1	0.82	0.59	0.72	0.04	-0.85
<mark>_0.0</mark>		0.19	0.001	0.04	0.05	0.03	0.02	0.06	0.03	0.04	0.01	0.92		0.29	0.87	0.09	0.29	0.17	0.96	0.07
II10rb RNAz 9.51 0.		0.59	0.8	0.16	0.4	0.42	0.74	0.5	0.79	0.77	0.67	-0.07	-0.17		-0.19		0.22	0.62		-0.72
	0	0.07	0.01	0.66	0.25	0.22	0.01	0.14	0.01	0.02	0.01	0.87	0.79	0.38	0.75	0.3		0.26	0.85	0.17
Cd14 RNAz 9.5 0.		0.55	0.84	0.78	0.44	0.43	0.66	0.57	0.65	0.52	0.54	-0.19			-0.06		-0.03			-0.63
	0	0.1	0.002	0.01	0.21	0.22	0.04	0.09	0.06	0.15	0.05	0.62	0.84	0.89		0.09	0.96	0.1	0.74	0.26
CTGF cells 9.28 0.		0.53	0.83	0.67		-0.46		0.61	0.7	0.3	0.7	0.31	-0.24		-0.94		-1			-0.93
0.0 C2 DNA- 0.10 0		0.28	0.04	0.15	0.72	0.36	0.11	0.2	0.12		0.04	0.55	0.85			0.46	0.05	0.93	0.55	0.25
Ccr2 RNAz 9.13 0.		0.41	0.91	0.48	0.5	0.44	0.5	0.3	0.76	0.68	0.61	-0.05 0.9	-0.38 0.53	0.16	-0.25 0.68	0.06	0.21	0.12	0.1	-0.7 0.19
Tgfb1 RNAz 9.12 0.		-0.01			0.15	0.49	0.15	0.4	0.02	0.04	0.02	-0.12		0.54	-0.3	0.61	0.74	0.63		-0.78
_	0.04	0.97	0.33	0.43	0.00	0.45	0.77	0.01	0.70	0.73	0.03	0.75		0.35	0.63	0.01	0.49	0.05	0.00	0.12
Tnfrsf1a RNAz 9.07 0.		-0.67		0.74	0.83	0.13	0.67	0.64	0.74	0.66	0.58	0.73		-0.36		0.61	0.45	0.23		-0.62
0.0		0.03	0.83		0.003		0.03	0.05	0.02	0.05	0.03	0.6	0.8	0.56	0.45	0.28	0.24	0.17	0.38	0.27
S100A4 cells 9.03 0.		0.2	0.54		0.69	0.04	0.83	0.9	0.96	0.2	0.29	0.73		-0.72		-0.69		0.97	0.84	-
0.		0.71	0.27	0.59	0.13	0.93	0.04		0.003		0.45	0.1		0.48	0.81			0.15	0.37	0.4
II1rn RNAz 8.99 0.	0.62	0.63	0.78	0.76	0.45	0.52	0.62	0.72	0.68	0.36	0.36	-0.32	0.32	-0.34	-0.08	0.87	0.59	0.84	-0.19	-0.77
0.0	01 0	0.05	0.01	0.01	0.2	0.12	0.06	0.02	0.05	0.34	0.21	0.4	0.6	0.58	0.9	0.06	0.29	0.08	0.81	0.13
Ifnar1 RNAz 8.92 0.	98 0.82	0.35	0.81	0.6	0.45	0.4	0.15	0.46	0.74	0.71	0.66	-0.01	0.5	0.06	-0.29	0.53	0.6	0.45	0.23	-0.61
	0	0.32	0.005	0.07	0.2	0.25	0.67	0.18	0.02	0.03	0.01	0.98	0.39	0.92	0.64	0.36	0.29	0.44	0.77	0.28
Bax RNAf 8.84 0.		0.76	0.82	0.28	0.53	0.46	0.64	0.62	0.72	0.65	0.27	-0.27	0.9	0.75	-0.02	0.34	0.49	0.74	0.35	-0.74
0.0			0.004		0.11	0.18	0.05	0.06			0.35	0.48				0.58	0.4	0.15	0.65	0.16
Slc10a1 RNAa 8.69 -0.			-0.61			-0.46	-0.7		-0.77					-			-0.62			-
0.		0.23	0.06	0.23	0.15	0.18	0.02	0.01	0.02		0.21	0.18	0.8			0.54	0.26	0.09	0.21	0.66
Smad6 RNAf 8.65 0.		0.01	0.5		0.27	0.49	0.61	0.53	0.78	0.73		-0.13	0.54	0.85	0.06		0.25	0.56		-0.49
DIII DEII- 0.00 0		0.98	0.14	0.05	0.45	0.15	0.06	0.12	0.01	0.03		0.74		0.07		0.11	0.68	0.33	0.84	0.41
BrdU BE cells 8.63 0.	0.86	0.5	0.2	-0.42 0.23	0.31	0.3	0.76	0.68	0.86	0.8	0.64	0.13	-0.53 0.36	-0.67 0.22	0.58	-0.59 0.29	0.55			0.33
Infarct area 8.61 0.		0.14	0.59	0.23	0.99	0.4	0.02	0.05	0.002	0.01	0.01	-0.01	0.30	0.22	0.56	0.29	0.55	0.27	0.71	0.59
	0.09	0.73	0.03		0.0002		0.39	0.08	0.32	0.42	0.41	0.98	n.a.	0.79	0.05		0.004	0.5	0.55	0.43
Hgf RNAz 8.26 0.		-0.39		-0.05		0.55	0.66	0.46	0.82	0.42	0.13	0.30	-0.03		-0.34		0.51	0.4		-0.35
0.20 0.		0.00	0.25	0.00	0.16	0.33	0.04	0.18	0.02	0.01	0.005		0.96	0.82	0.57	0.02	0.38	0.58	0.49	0.57
Ccl7 RNAz 8.21 0.		-																		-0.6
	96 0.79	0.39	0.56	0.35	0.6	0.33	0.2	0.43	0.63	0.63	0.6	-0.05	-0.05	-0.48	-0.23	0.17	0.55	0.6	0.02	-0.0

4.5 BrdU Hep cells

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B 0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
Mki67 RNAf	10.4	0.9	0.74	0.91	0.65	-0.5	0.57	0.44	0.86	0.87	0.91	0.49	0.55	0.82	0.94	0.94	-0.08	-0.57	0.98	0.81	0.9	0.89
		0.002	0	0.0002	0.04	0.14	0.09	0.21	0.001	0.001	0.0003	0.15	0.03	0.003	0.02	0.02	0.9	0.31	0.002	0.09	0.03	0.05
Birc5 RNAf	10.3	0.87	0.72	0.78	0.69	-0.57	0.61	0.45	0.9	0.92	0.9	0.47	0.53	0.84	0.23	0.83	-0.25	-0.88	0.81	0.89	0.9	0.93
		0.005	0	0.01	0.03	0.09	0.06	0.2	0.0003	0.0002	20.0003	0.17	0.04	0.003	0.71	0.08	0.68	0.05	0.1	0.04	0.03	0.02
Notch1 RNAf	8.79	0.76	0.76	0.51	0.16	-0.55	0.42	0.07	0.92	0.93	0.84	0.82	0.63	0.44	-0.4	0.59	-0.03	-0.67	0.44	0.96	0.82	0.76
		0.03	0	0.13	0.65	0.1	0.23	0.84	0.0001	0	0.003	0.003	0.01	0.21	0.51	0.3	0.96	0.22	0.46	0.01	0.09	0.13
Lama1 RNAf	7.71	0.67	0.57	0.81	0.79	-0.24	0.01	0.08	0.75	0.57	0.88	0.38	0.32	0.64	-0.77	0.91	-0.53	-0.3	0.02	0.61	0.82	0.73
		0.07	0.0001	0.005	0.01	0.51	0.98	0.82	0.01	0.08	0.001	0.27	0.24	0.05	0.13	0.03	0.36	0.63	0.98	0.27	0.09	0.16
BrdU NP cells	7.67	0.88	0.62	0.53	0.56	0.5	0.24	0.27	0.55	0.49	0.66	0.19	0.31	0.39	-0.27	0.42	-0.17	0.92	0.17	0.19	-0.22	0.91
		0.004	0	0.12	0.09	0.14	0.51	0.46	0.1	0.15	0.03	0.58	0.25	0.23	0.66	0.48		0.03		0.76	0.68	0.03
Cdh2 RNAf	7.37	0.6	0.72	0.11	-0.26	-0.47		-0.13	0.86	0.89	0.61	0.84	0.76	0.33	-0.03	0.71	-0.34	-0.79	0.87	0.89	0.81	0.18
		0.12	0		0.48	0.17					0.06				0.96	0.18	0.58			0.04	0.1	0.77
Nes RNAf	7.27	0.74	0.65		-0.23		0.4	0.08		0.83	0.82		0.44	0.45		-0.13		-0.66		0.91	0.81	0.5
		0.03	0	0.65		0.15					0.003		0.1	0.2				0.23		0.03	0.1	0.39
Mrc1 RNAz	7.15	0.55	0.39	0.63		-0.24		0.6		0.83				-0.5				-0.28	0.82	0.9		-0.34
		0.16	0.01	0.05	0.05	0.5	0.03				0.003			0.14		0.18		0.65		0.04		0.57
Por RNAa	6.88	-0.51	-0.41	0.09		-0.58		-0.4			-0.71				0.2							-0.9
		0.2	0.01	0.81	0.76	0.08		0.25	0.22	0.06		0.14		0.004	0.75	0.65		0.23		0.23	0.89	0.03
Timp2 RNAf	6.74		0.4			-0.63			0.89	0.86			-0.06		0.69			-0.62		0.9		-0.28
		0.23	0.01								0			0.32	0.2	0.05		0.27		0.04	0.03	
Cdh1 RNAf	6.57	0.64	0.64			-0.55						0.7			-0.48			-0.66		0.8	0.96	0.81
		0.09	0	-	0.36		0.49	0.99	0.01			0.02		0.07	0.41	0.9	0.88	-	0.77	0.1	0.01	0.1
CTGF cells	6.34	0.59	0.42		-0.45		0.72	0.78	0.75	0.73							-1	0.98	0.78	0.09	0.83	0.78
			0.04	0.3		0.47		0.07	0.09		0.001		0.16		0.72		0.06	0.14			0.38	
Slc10a1 RNAa	6.2	-	-0.49	0.12							-0.85						-0.92		0.55		-0.49	
6	0.40		0.001		0.53	0.01		0.69			0.002			0.003		0.66		0.04			0.4	0.06
Gstm1 RNAa	6.18	-	0.57		-0.33			-0.07		0.82			0.44					-0.85		0.8	0.46	-0.6
A . I . D. N. A	0.00		0.0001	-		0.02	0.67	0.84		0.004		0.01	0.1	0.46	0.5	0.9		0.07		0.11	-	0.29
Actb RNAz	6.08	0.65			-0.37					0.59	0.74			0.54	-0.43		0.75		0.04	0.63	0.95	0.64
C DNIAC	0.04		0.0002				0.55	0.93	0.17		0.02		0.11	0.1	0.47		0.14			0.25	0.01	0.25
Sparc RNAf	6.01	0.58						-0.02	-		0.94				-0.58						0.93	0.28
C IC DNAC	F 0.4		0.003	0.05	0.2		0.91	0.95	0.01	0.04	0		0.97	0.76			0.49		0.97		0.02	
Smad6 RNAf	5.94	0.63			0.22	0.08	0.14	0.11	0.42	0.58		0.38	0.11	0.23	-0.26			0.13		0.64	0.82	
D III IV "	F 00		0.001	0.1	0.55	0.83	0.69	0.75	0.23		0.001		0.7	0.53	0.67		0.86	0.84		0.24	0.09	
BrdU Kup cells	5.69		0.52	0.55		0.15		0.04	0.6		0.14			0.13	-0.36			-0.29	-0.1	0.4	-0.83	
		0.01	0.001	0.1	0.09	0.68	0.92	0.91	0.08	0.17	0.69	8.0	0.45	0.7	0.55	0.41	0.21	0.63	0.87	0.6	0.04	0.01

4.6 BrdU NP cells

4.6 BrdU NP cells

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2	:B0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
BrdU Kup cells	11.8	0.87	0.82	0.81	0.89	0.47	0.96	0.96	0.96	0.92	0.68	0.28	0.61	0.85	-0.28	0.89	0.32	0.06	0.94	0.89	0.17	0.97
		0.005	0	0.005	0.001	0.17	0	0	0	0.0004	4 0.02	0.43	0.02	0.001	0.64	0.04	0.6	0.93	0.02	0.11	0.75	0.01
BrdU BE cells	7	0.95	0.89		-0.05	0.6	0.39	0.74	0.9	0.83	0.9	0.71	0.74	0.74	-0.15	-0.41	0.75	-0.28	0.51	0.71	-0.78	0.84
		0.0003		0.38	0.88	0.07	0.26	0.02	0.001	0.01	0.0004	0.03	0.002	0.01		0.49		0.64	0.38	0.29	0.12	
Mki67 RNAf	9.49		0.79	0.57	0.35	-0.53	0.62	0.61	0.84	0.75	0.7	0.5	0.54	0.45	-0.43	0.4		-0.83		0.61	-0.08	-
		0	0	0.08	0.33	0.12	0.06		0.002		0.03	0.14	0.04	0.19	0.47	0.51		0.08	0.9	0.27	0.9	0.18
Birc5 RNAf	9.1	0.99	0.82	0.33	0.46	-0.14		0.4	0.83	0.69	0.71	0.52	0.64	0.6	0.06	0.29		-0.98		0.56	0.06	0.83
C . 0 DNA	0.00	0	0	0.36	0.18	0.71	0.38		0.003		0.02	0.12	0.01	0.06		0.63		0.003		0.33	0.92	0.08
Gsta2 RNAa	8.06		0.6		-0.13		0.83	0.65	0.9	0.94	0.86	0.69	-0.13			-0.76		-0.93		0.89		-0.56
1 1 DNAC	0.04	0.001	0 00	0.68	0.71				0.0004	-		0.03	0.64	0.16	0.32		0.5	0.02	0.08	0.04	0.37	
Lama1 RNAf	8.01		0.66	0.53	0.67				0.84		0.66	0.54	0.46	0.28	-0.18		0.11	-0.56			-0.42	= =
Callal DNAS	7.89	0.005	0	0.11	0.04	-		0.71	0.002	0.1	0.04	0.11	0.09	0.43	0.77	0.92	0.86	0.33		0.17	0.48	0.42
Col1a1 RNAf	7.09	0.002	0.78	-	0.47	-0.61 0.06	0.29		0.02		0.71	0.51	0.5	0.41	0.71	0.44		-0.89 0.04	0.59	0.47	0.01	0.72
CTGF cells	7.88	0.00_	0.77	0	•		• • • •	0.49	0.86	0.00	0.02			0.12	1			0.04			0.62	0.17
CTGI Cells	7.00	0.004		0.05	0.01	0.34	0.07	0.61	0.00		0.004		0.23	0.12		0.20		0.34			0.02	
Mrc1 RNAz	7 77	0.87		0.72	0.83		0.09	0.31	0.75					-0.38				-0.62				-0.39
WITCH TATAL	1.11	0.005	0.02		0.003	0.72	0.81	0.39	0.01		0.004			0.27	0.97		0.77			0.65	0.75	
BrdU Hep cells	7.67		0.62		0.56	0.5	0.24	0.27	0.55	0.49			0.31	0.39				0.92			-0.22	
2.40 .10p co		0.004			0.09	0.14		0.46	0.1	0.15			0.25	0.23	0.66			0.03		0.76	0.68	
S100A4 cells	7.62	0.91	0.79	-	-0.96			0.46	0.94	0.87	0.87			0.08	0.97		1	-0.92			-0.38	
		0.002			0.002			0.36	0.01	0.02				0.88	0.14		0.03			0.59		0.72
Pde4a RNAf	7.44	0.89	0.73	-0.43	-0.52	-0.33	0.44	0.48	0.69	0.35	0.58	0.62	0.58	0.45	-0.28	0.33	-0.45	-0.5	0.15	0.8	-0.13	0.72
		0.003	0	0.21	0.12	0.35	0.2	0.16	0.03	0.32	0.08	0.06	0.02	0.2	0.65	0.59	0.45	0.39	0.81	0.1	0.83	0.17
Slc10a1 RNAa	7.3	-0.81	-0.63	0.45	0.56	-0.04	-0.16	-0.04	-0.58	-0.58	-0.82	-0.72	-0.71	-0.74	0.51	0.84	0.25	-0.87	0.23	-0.64	0.37	-0.92
		0.01	0	0.19	0.09	0.9	0.65	0.91	0.08	0.08	0.003	0.02	0.003	0.01	0.38	0.08	0.69	0.05	0.71	0.24	0.54	0.03
Ugt1a1 RNAa	7.23	-0.77	-0.64	0.39	0.14	-0.67	0.3	-0.02	-0.21	-0.5	-0.9	-0.73	-0.7	-0.68	0.42	-0.23	-0.15	-0.98	-0.32	-0.67	0.33	-0.8
		0.03	0	0.26	0.69	0.03	0.4	0.95	0.57		0.0003				0.48			0.003	0.6	0.21	0.58	0.1
Tnfrsf1a RNAz	7.08	0.82		0.6		-0.16		0.32	0.55		0.53				-0.64		-0.92			0.72		
			0.0001	0.06	0.34	0.66	0.03	0.36	0.1	0.06	0.12				0.24		0.03	0.26	-	0.17	0.45	
Sparc RNAf	7	0.89	0.65	0.51	0.42	-0.32		0.07	0.83	0.53	0.75		0.22			-0.14				0.64		
C 0 0= D:::	0.00	0.003	0	0.13	0.23	0.36	0.99		0.003		0.01	0.07			0.08	0.83	0.93	0.1		0.25	0.89	0.98
Cyp2c37 RNAa	6.96		-0.65	0.67	0.4	-0.39		-0.05		-0.24		-0.7	-0.55		-0.22		0.1	-0.1		-0.57		
C-+1 DNA	6.00	0.01	0	0.03	0.25	0.26	0.06	0.88	0.04	0.5	0.01	0.02	0.03	0.1	0.72	0.93	0.87	0.87	0.1	0.31	0.9	0.11
Gstm1 RNAa	6.88	0.82	0.63	-		-0.79		0.5	0.8	0.61	0.62		0.07	-0.1		-0.05	0.06		-0.77	0.59		-0.24
		0.01	0	0.22	0.12	0.01	0.06	0.14	0.01	0.06	0.05	0.33	0.82	0.77	0.75	0.94	0.92	0.04	0.13	0.3	0.04	0.7

4.7 BrdU Kupffer cells

Factor	Cons	Av	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	30h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
BrdU NP cells	11.8	0.87	0.82	0.81	0.89	0.47	0.96	0.96	0.96	0.92	0.68	0.28	0.61	0.85	-0.28	0.89	0.32	0.06	0.94	0.89	0.17	0.97
		0.005	0	0.005	0.001	0.17	0	0	0	0.0004	0.02	0.43	0.02	0.001	0.64	0.04	0.6	0.93	0.02	0.11	0.75	0.01
BrdU BE cells	7.45	0.69	0.67	-0.27	0.11	0.33	0.3	0.65	0.94	0.81	0.58	-0.001	0.27	0.74	0.24	-0.12	-0.04	0.4	0.4	0.82	0.28	0.84
		0.06	0	0.45	0.77	0.35	0.39	0.04	0.0002	0.01	0.08	1	0.35	0.01	0.7	0.84	0.96	0.51	0.51	0.18	0.64	0.07
Mki67 RNAf	7.11	0.84	0.63	0.47	0.37	-0.07	0.45	0.46	0.91	0.73	0.27	-0.08	0.22	0.28	-0.03	0.31	0.45	-0.45	-0.22	0.8	-0.85	0.78
		0.01	0	0.17	0.3	0.85	0.19	0.18	0.001	0.03	0.45	0.84	0.44	0.44	0.96	0.61	0.44	0.45	0.72	0.2	0.07	0.12
Gsta2 RNAa	6.79	0.81	0.52	0.04	0.02	-0.33	0.77	0.65	0.9	0.95	0.51	0.27	-0.33	-0.61	-0.17	-0.38	-0.68	-0.28	0.82	0.99	-0.61	-0.53
		0.01	0.001	0.92	0.96	0.35	0.01	0.04	0.001	0	0.13	0.48	0.25	0.06	0.78	0.53	0.21	0.65	0.09	0.01	0.28	0.36
Gstm1 RNAa	6.71	0.93	0.69	0.21	-0.27	-0.51	0.53	0.45	0.83	0.6	0.07	0.59	0.2	-0.35	-0.3	0.24	-0.83	-0.21	-0.9	0.79	-0.41	-0.35
		0.001	0	0.57	0.45	0.14	0.12	0.19	0.01	0.09	0.86	0.09	0.49	0.33	0.63	0.7	0.08	0.73	0.04	0.21	0.49	0.57
S100A4 cells	6.67	0.93	0.78	-0.46	-0.67		0.43	0.22	0.92	0.72	0.53		0.01	-0.02	-0.89	-0.5	0.97	0.8	-0.86	1	0.16	0.27
		0.001	0	0.35	0.14	0.71	0.4	0.68	0.03	0.17	0.28	0.71	0.98	0.97	0.3	0.67	0.15	0.41	0.34	0	0.9	0.83
Birc5 RNAf	6.56	0.8	0.61	0.21	0.41	0.33	0.13	0.23	0.89	0.63	0.24	-0.13	0.26	0.39	0.18	0.12	0.12	-0.18	-0.64	0.75	-0.87	0.87
		0.02	0	0.56	0.24	0.36	0.71			0.07		0.74			0.77	0.85	0.85			0.25	0.06	0.05
Rarres1 RNAf	6.38	-0.81	-0.49	0.25	0.24	0.64	-0.32	-0.65	-0.85	-0.71	-0.25	-0.57	-0.42	0.06	0.65	-0.41	8.0	-0.22	-0.52	-0.79	0.03	0.37
		0.02		0.49	0.5	0.04	0.37	0.04	0.003	0.03	0.48	0.11	0.14	0.87	0.23		0.1		0.37		0.96	0.53
Slc10a1 RNAa	6.18	-0.77	-0.61	0.3				0.001		-0.55					0.03		-0.36			-0.76		-0.9
		0.02	0	0.39	0.02		0.6	1	0.12	0.13	0.1	0.38	0.1	0.02		0.03	0.55	0.84		0.24		0.04
CTGF cells	6.08	0.74	0.63			-0.47			0.99	0.91		-0.68			-0.76	0		-0.84		-1	-0.42	
		0.04		-						0.03		0.21			0.45	1			0.82	0	0.72	
GLDH blood	5.99	-0.41								-0.48			0.28				-0.94		-0.59		-0.06	
			0.04			0.04		0.11		0.19			0.33		0.43		0.02		0.29	0.82	0.92	
Por RNAa	5.72	-0.07				-0.18				-0.79			-0.77		0.12					-0.92		
		0.87	0.02	0.58	0.84	0.61	0.49	0.79	0.38			0.08			0.84		0.6	0.39	0.16		0.67	
BrdU Hep cells	5.69		0.52	0.55		0.15		0.04	0.6	0.5			0.21		-0.36			-0.29	-0.1		-0.83	
	5 40	0.01		0.1	0.09			0.91	0.08		0.69	0.8	0.45	0.7		0.41		0.63	0.87		0.04	
Mrc1 RNAz	5.48	0.59		0.57		0.14				0.43	0.52		-0.28							0.63		
T (() DNA	- 00		0.003	0.09	0.01	0.7			0.003		0.12		0.34	0.32	0.91	0.2	0.68			0.37		
Tnfrsf1a RNAz	5.38	0.68				-0.5	0.64		0.51	0.58	0.39	0.22		-0.2						0.72		
C 4 10 DNA	F 00	0.06		0.35	0.91	0.14	0.05		0.16	0.1	0.26	0.57	0.96	0.57	0.81	0.84	0.8			0.28		0.8
Cyp4a10 RNAa	5.03			0.45	0.15					-0.53		-0.69			-0.38	0.1	0.61		-0.005			-0.34
I 1 DNAC	4.0		0.01	0.19	0.68	0.91			0.14		0.41	0.04		0.81	0.53	0.87	0.28					
Lama1 RNAf	4.9	0.58	0.41	0.39	0.6				0.93				0.04	0.2	0.05	0.15	0.08		-0.55		-0.86	
D 11 DNAC	4.07		0.01	0.26	0.07	0.34	0.5		0.0002		0.52	0.69	0.89	0.58	0.93	0.8	0.9	0.51		0.11	0.06	-
Bak1 RNAf	4.87	0.72	0.6		-0.53		0.03	0.11		0.44			0.27	0.32		-0.59			-0.52		-0.91	
		0.05	0	0.3	0.11	8.0	0.93	0.77	0.01	0.24	0.84	0.69	0.35	0.36	0.21	0.29	0.22	0.08	0.37	0.02	0.03	0.08

4.8 BrdU+ biliary epithelial cells

Factor	Cons	Av	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B 0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
BrdU NP cells	10.2	0.95	0.89	-0.31	-0.05	0.6	0.39	0.74	0.9	0.83	0.9	0.71	0.74	0.74	-0.15	-0.41	0.75	-0.28	0.51	0.71	-0.78	0.84
		0.0003	0	0.38	0.88	0.07	0.26	0.02	0.001	0.01	0.0004	0.03	0.002	0.01	0.81	0.49	0.14	0.64	0.38	0.29	0.12	0.07
S100A4 cells	9.3	0.82	0.86	0.82	0.35	-0.26	0.37	0.77	0.99	0.98	0.91	0.12	0.27	0.36	-0.49	1	0.88	0.11	0.54	1	0.25	0.75
		0.01	0	0.05	0.49	0.62	0.47				0.01		0.51	0.49	0.67	0	0.31		0.64	0	0.84	
Col1a1 RNAf	8.69	0.97	0.85	-	-0.04	-0.12			0.98		0.61	0.46	0.64	0.35		-0.54	-0.003		0.58	0.98	-0.72	0.78
		0	0	0.55	0.92	0.74	0.27	0.02		0.0004		0.25	0.02	0.36	-	0.35	1	0.95	0.3	0.02	0.28	0.12
Bili blood	8.63		0.86			-0.42		0.3	0.76		0.86	0.8	0.64			-0.67		-0.59	0.36		-0.23	
T (14 DNA	0.00	0	0	0.14	0.2	0.23	0.39	0.4	0.02		0.002		0.01	0.71	0.36	0.22	0.58	0.29	0.55	0.27	0.71	0.59
Tgfb1 RNAz		0.95	0.78	-	-0.27	-0.4	0.27	0.81	0.9	0.7	0.9		0.47			-0.94		0.02	0.72	0.85		-0.42
U10 L DNIA		0.0003		0.23	0.45	0.26					0.001		-	0.64	0.39	0.02	0.39	0.98	0.17	0.15	0.39	
II10rb RNAz		0.95		0.1		-0.22	0.2	0.69	0.82				0.49		-0.39		-0.6	0.14	0.69	0.91		-0.41
C 2 - 27 DNA -		0.0004	_	0.78	0.81	0.55	0.58	0.03	0.01		0.002				0.52		0.29	0.83	0.19	0.09	0.52	0.5
Cyp2c37 RNAa	0.09	0.004	-0.7 0	-0.47		-0.45 0.2	0.84	-0.4 0.25	-0.53		-0.82 0.01										-0.02	
Slc10a1 RNAa	8.04	-0.8		0.17	0.9	0.2		-0.29			-0.88					0.22		0.47			0.98	0.23
SICIUAL KINAA	0.04	0.02	0.00	0.52	0.09	0.04	0.4	0.41	0.05		0.002			0.2	0.24		0.79	0.07		0.01		0.3
Ugt1a1 RNAa	7.9		-0.67		0.46			• • • • •			-0.95			-	-0.73		-0.6				-0.47	
Ogital MNA	1.0	0.03	0.07	0.09	0.18	0.54	0.18	0.10	0.47				0.70			0.28	0.28	0.62	0.02		0.53	0.30
Cyp2e1 RNAf	7.86	-0.93	•	-0.41	-0.2	-0.59		0.09	-0.6		-0.88					0.13		-0.4			-0.27	
Cypzei Kit/Ki	7.00	0.001	0.72	0.25	0.59	0.07	0.85	0.81	0.09		0.002		0.03	0.82	0.03		0.49	0.51	0.53	0.43	0.73	0.76
Ccr2 RNAz	7.69	0.92	0.75		-0.31		0.3	0.69	0.67	0.58	0.88	0.82		-0.2		-0.72						-0.44
00.2	7.00	0.001	0	0.97	0.38	0.24	0.39	0.03	0.05	0.1	0.002			0.6	0.88	0.17	0.36		0.21	0.58		0.45
CTGF cells	7.64	0.9	0.89	0.47	0.16		-0.18	-0.48		0.85	0.81	0.31	0.52	0.02		-0.87		-0.19		-1	-0.51	
	-	0.002	0	0.35		0.12	0.73	0.33	0.03	0.07	0.05	0.61	0.18	0.96		0.33	0.66	0.88	0.15	0		0.31
Hgf RNAz	7.57	0.95	0.71	-0.46	0.02	-0.06	0.36	0.72	0.8	0.31	0.83	0.82	0.38	-0.3	-0.4	-0.005	-0.38	-0.27	0.7	0.67	0.61	-0.69
· ·		0.0003	0	0.19	0.95	0.87	0.3	0.02	0.01	0.41	0.01	0.01	0.2	0.44	0.51	0.99	0.52	0.67	0.19	0.33	0.39	0.2
Cd86 RNAz	7.49	0.95	0.8	-0.45	-0.21	-0.46	-0.22	0.46	0.75	0.51	0.88	0.86	0.53	-0.04	-0.33	-0.48	-0.67	0.02	-0.22	0.59	0.61	-0.21
		0.0003	0	0.19	0.57	0.18	0.55	0.18	0.02	0.16	0.002	0.01	0.06	0.92	0.59	0.41	0.22	0.97	0.73	0.41	0.39	0.74
Lama1 RNAf	7.48	0.9	0.68	-0.53	-0.09	0.09	0.27	0.67	0.85	0.4	0.72	0.66	0.53	0.11	-0.9	-0.54	-0.4	-0.26	0.55	0.68	-0.44	0.27
		0.002	0	0.12	0.8	0.81	0.45	0.03	0.004	0.29	0.03	0.07	0.06	0.77	0.04	0.35	0.5	0.67	0.34	0.32	0.56	0.66
BrdU Kup cells	7.45	0.69	0.67	-0.27	0.11	0.33	0.3	0.65	0.94	0.81	0.58	-0.001	0.27	0.74	0.24	-0.12	-0.04	0.4	0.4	0.82	0.28	0.84
		0.06	0	0.45	0.77	0.35	0.39	0.04	0.0002	0.01	0.08	1	0.35	0.01	0.7	0.84	0.96	0.51	0.51	0.18	0.64	0.07
Ccl2 RNAz	7.44		0.75		-0.16	-0.38	0.28	0.47	0.64	0.61	0.82	0.82	0.44	-0.18			-0.48		0.68	0.89	0.47	
		0.002	0	0.89	0.66	0.28	0.43	0.17	0.06	0.08	0.01	0.01	0.13	0.65	-	0.82	0.41	0.87	0.21	0.11		0.49
Sparc RNAf	7.36		0.7			-0.01	0.26	0.59	0.87	0.67	0.71	0.61	0.32	-0.36		-0.43			0.43		-0.67	
		0.0002	0	0.07	0.99	0.97	0.48	0.08	0.003	0.05	0.03	0.11	0.28	0.34	0.41	0.47	0.52	0.97	0.47	0.22	0.33	0.64

4.9 Collagen area

Factor	Cons		All							30h-2d							T12h			T2d		T14d
Tnfrsf1b RNAz	7.51	0.92	0.78	0.16	0.52	0.7	0.28	0.99	0.67	0.01	0.46		0.84		-0.58 0.31		-0.71 0.18	0.59	-0.48 0.41	0.79	0.59	0.35
Osm RNAz	7.39	0.94	0.7	-0.1		-0.29				0.45	0.55	0.71	0.66		-0.42		-0.76		0.87	0.75	-	-0.37
		0.001	0	0.77	0.18	0.42	0.98	0.45	0.06	0.2	0.13	0.03	0.01	0.26	0.48	0.32	0.13	0.42	0.06	0.14	0.39	0.54
Ccr5 RNAz		0.95	0.77	-0.27	0.6	-0.32			0.08	-0.13	0.6	0.77	0.78	0.62	-0.34		-0.83		-0.26			-0.08
Osmr RNAz		0.0004 0.91	0.73	0.44	0.06	0.37	0.23	0.55	0.82	0.71	0.09	0.01	0.001		0.57	-0.6	0.08	0.04	0.67 -0.38	0.79	0.29	0.9 -0.25
OSIIII KINAZ	- 1	0.002		0.28	0.13	0.2	0.32	0.84		0.81	0.14		0.73		0.61	0.28		0.92	0.53	0.97	0.7	0.68
Ccl2 RNAz	6.74	0.92		0.001			-0.3	0.04	0.66	0.06	0.22	0.48	0.78	0.6			-0.79		-0.22		0.13	0.35
		0.001	0	1	0.09	0.62	0.4	0.9	0.04	0.88	0.58	0.19	0.001	0.09	0.23	0.17	0.12	0.27	0.73	0.11	0.87	0.56
Cd86 RNAz	6.68	0.89	0.74	-0.29		-0.15			0.31	0.03	0.4	0.6	0.74		-0.57		-0.6	0.83	-0.05		0.42	0.04
II10	6.60	0.003	0 71	0.41	0.18	0.68	0.66	0.31	0.39	0.93	0.29		0.002		0.31	0.9	0.28	0.08		0.48	0.58	0.95
II10rb RNAz	6.63	0.9	0.71	0.22	0.61	-0.08 0.82	0.29	0.07	0.54	-0.01 0.98	0.29	0.55	0.71		-0.41 0.5	0.31	-0.74 0.15	0.63	-0.31 0.62		0.26	-0.17 0.79
II2 RNAz	6.57	0.95	0.68	0.34	0.13			-0.22		0.13	0.6	0.76	0.64		-0.5	0.77			0.004		-	-0.38
		0.0004	0	0.34	0.72	0.68	0.68	0.55	0.69	0.72	0.09	0.02	0.01	0.26	0.39	0.13		0.17	0.99	0.36	0.31	0.53
Tnfrsf1a RNAz	6.55	0.61	0.55	-0.74			-0.09	0.29	0.85		0.61	0.76	0.67	0.51			-0.54		-0.08	0.96	0.74	-0.01
M. DNIA	0.54		0.0002		0.39	0.97	0.81		0.002		0.08	0.02	0.01	0.16	0.58	0.37		0.05	0.89	0.01	0.26	0.98
Met RNAz	6.54	7 7	0.56 0.0002	0.56	0.29	0.56	0.22	0.15	0.07	-0.07 0.84	0.52	0.6	0.52	0.3	0.35	-0.54 0.35	0.57	0.52	-0.01 0.99	-0.43 0.47	0.61	-0.51 0.38
Bili blood	6.51	0.01	0.64	0.09		-0.27				-0.14	0.13	0.09	0.64	0.43	-0.77	0.33	-0.5	0.37	-0.28	0.47		-0.44
Dill blood	0.01	0.02	0.01	0.12	0.1			0.91	0.27	0.7	0.2	0.01	0.01	0.34	0.13	0.01	0.39	0.62		0.08	0.05	0.45
II13 RNAz	6.37	0.93	0.69	-0.29	0.38	-0.5	-0.42	-0.16	0.18	0.13	0.6	0.78	0.65	0.4	-0.08	0.05	-0.9	0.21	0.004	0.53	0.74	-0.35
		0.001	0	0.41	0.29			0.65	0.61	0.72	0.09	0.01	0.01	0.28	0.89	0.93	0.04		0.99	0.36		0.56
Cxcl3 RNAz	6.33	0.92	0.68	-0.13	0.29	-0.37				0.14	0.6	0.77	0.65	0.4	-0.5	-0.61	-0.86		0.04	0.5		-0.39
Ccr2 RNAz	6 20	0.001	0 0.74	0.73	0.42	0.3	0.3	0.75	0.45	0.71	0.09	0.02	0.01	0.28	0.39	0.28	0.06		0.95 -0.13	0.39	0.3	0.51
CCIZ KNAZ	0.02	0.001	0.74	0.03	0.08		0.61	0.14	0.13	0.75	0.41		0.002		0.78	0.01	0.21	0.01		0.20	0.20	0.1
Ifna1 RNAz	6.31	0.95	0.71	-0.13	0.05	-0.58		-0.3		-0.06	0.79	0.93	0.7	0.53	0.41				-0.23		• · · ·	-0.77
		<mark>0.000</mark> 4	0	0.72	0.9	0.08	0.45	0.39	0.3	0.86	0.01	0.0003	0.01	0.15	0.49	0.21	0.32	0.28	0.7	0.76	0.05	0.13
Cxcr1 RNAz	6.28	7 7	0.75			-0.69			-0.27		0.7	0.86	0.75	0.57	0.26				-0.16		0.84	
D DNAC		0.0005		0.42	1		0.34	0.46	0.45	0.97			0.002		0.67	0.55		0.38				0.54
Pparg RNAf	b.14	0.89	0.6	0.77	0.2 0.58	0.13	0.31	0.13	-0.07 0.85	0.16	0.2	0.13	0.61		-0.28 0.65	0.42	0.61	-0.36	0.2	0.25		0.15
Cyp3a11 RNAa	5.75		-0.26			-0.68		-0.31					-0.68						-0.18		-0.27	-0.4
0,,00011 1111/10	3.70	0.55	0.11	0.01		0.03				0.47			0.01		0.04				0.77		0.73	0.5

4.10 Infarct area 15

4.10 Infarct area

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
Bili blood	8.61	0.98	0.69	0.75	0.59	0.41	0.92	0.6	0.39	0.58	0.32	0.29	0.41	-0.01	0	0.79	0.88	0.11	0.98	0.5	-0.36	0.43
		0	0	0.01	0.07	0.24	0.0002	0.07	0.27	0.08	0.37	0.42	0.13	0.98	n.a.	0.11	0.05	0.86	0.004	0.4	0.55	0.47
II17a RNAz	8.25	0.92	0.81	-0.07	0.43	0.05	0.15	0.02	0.44	0.5	0.84	0.82	0.78	0.77	0	-0.24	-0.34	-0.03	0.16	0.64	0.89	0.46
		0.001	0	0.86	0.22	0.89	0.69	0.96	0.2	0.14	0.005	0.01	0.001	0.01	n.a.	0.7	0.58	0.96	0.79	0.25	0.11	0.43
SMA cells	7.76	0.96	0.73	0.6	0.74	0.21	-0.45	-0.52	0.44	-0.27	0.57	0.67	0.59	0.56	0	-0.94	-0.82	0.36	-0.99	0.08	0.84	0.83
		0.0001	-	0.21	0.09	0.69	0.37		0.39	0.6	0.24	0.14	0.09	0.25	n.a.	0.21					0.37	0.37
Actb RNAa	7.71		0.66	0.44	0.31	0.28	0.55	-0.21	0.13	0.22	0.86	0.73	0.5	0.5	0	0.76	0.62	-0.56	0.21	0.26	0.88	-0.43
		0.001	0	0.2	0.38	0.44	0.1	0.55	0.71		0.003		0.07	0.17	n.a.	0.13		0.32			0.12	-
Actb RNAf	7.64		0.62	0.62	0.4	0.18		-0.31		0.27	0.9	0.78	0.48	0.48	0		0.53					-0.67
		0.005	0	0.06	0.25	0.61	0.21		0.75			0.01	0.08	0.2	n.a.		0.36				0.06	
Cyp1a2 RNAa	7.61		-0.58		-0.72						-0.63		-0.34	-0.6	0	-0.46						0.002
			0.0001	0.15	0.02	0.1	0.04	0.1	0.96	0.69	0.07		0.23	0.09	n.a.				0.27		0.23	1
Actb RNAz	7.39		0.69	0.36	0.32		0.28	-0.3	0.06	0.3		0.83	0.58	0.54	0	-0.18	-	-0.71		0.36		-0.69
		0.001	0	0.3	0.37		0.43	0.4	0.86		0.002		0.03	0.13	n.a.	0.77	0.87	0.18	0.72	0.55	0.08	
CTGF cells		0.95	0.71	0.45				-0.69		0.2			0.41	0.25	0	0	-1	-0.01		-1		-0.87
		0.0003	-		0.12			0.13	0.47	0.71	0.07			0.63	n.a.	1	0.02		0.14			0.33
Notch3 RNAf	7.22		0.7		-0.64					0.44		0.88	0.64	0.51	0				0.27			-0.42
		0.003	-	0.004		0.07		0.82	0.5		0.002			0.16	n.a.	0.09				0.11	0.03	0.48
Col3a1 RNAf	7.16		0.7		-0.42		0.5	0.15	0.23	0.34	0.85			0.42	0		-0.42			0.38		-0.26
		0.001	0		0.22		0.14			0.34				0.26	n.a.	0.3	0.49		0.6		0.04	
Pde4a RNAf	7.06		0.63	-	0.37			0.1	0.31	0.28	0.85			0.3	0		-0.45			0.4		-0.86
51.151116		0.001	0	0.7	0.3	0.31	0.46	0.77	0.39		0.003			0.44	n.a.	0.98	0.44		0.96	0.5	0.01	0.06
Pde4d RNAf	7.06		0.59	0.29	0.2	0.39	0.04	-0.1		0.22	0.74		0.57	0.45	0	0.67	0.76		-0.41			-0.27
Haal Dala		0.001	0	0.42	0.58	0.27		0.79		0.55	0.02	0.01	0.03	0.23	n.a.	0.21	0.13		0.49	0.38	0.19	0.66
II28b RNAz	- /	0.9	0.69		0.43		0.15	0.02	0.44	0.5	0.61	0.6	0.58	0.43	0	-0.24				0.64	0.58	0.34
C II I DNIAC	0.04	0.002	0	0.86	0.22	0.89	0.69	0.96	0.2	0.14	0.08	0.08	0.03	0.24	n.a.	0.7	0.58		0.79	0.25	0.42	0.58
Col1a1 RNAf	6.94		0.72		-0.34		0.5	0.12		0.13			0.58	0.44	0		-0.38			0.17	0.93	-0.3
CL 10 1 DNA		0.0003		-			0.15		0.66	0.72	0.01			0.23	n.a.	0.8	0.53		0.71		0.07	
Slc10a1 RNAa	6.93		-0.57		-0.47					-0.45				0.1	0				-0.56		-0.82	
D 1 DNAC	0.00		0.0002				0.22		0.26	0.19	0.11		0.35	0.79	n.a.	0.77			0.32			0.07
Prom1 RNAf		0.95			-0.15						0.9	0.92	0.7	0.63	0				-0.61			-0.45
D DNAC		0.0003		0.86		0.28	0.86	0.18					0.005		n.a.	0.62	0.73				0.04	
Bax RNAf	6.9	0.94	0.62	0.35	0.29	-0.5		-0.07		0.49	0.73		0.39	0.2	0		-0.33			0.54		-0.75
II .1 1 DNA	C 7E	0.001	0	0.33	0.41	0.14	0.3	0.84			0.02	0.03	0.17	0.6	n.a.	0.47	0.59		0.49		0.27	
Ugt1a1 RNAa	0./5		-0.41		-0.51					-0.77				0.3	0				-0.67			
		0.02	0.01	0.18	0.13	0.55	0.87	0.13	0.03	0.01	0.18	0.14	0.61	0.43	n.a.	0.49	0.58	0.31	0.22	0.08	0.59	0.03

4.11 SMA cells

Factor	Cons	Av	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2c	B0h-5d	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
CTGF cells	11.9	0.98	0.93	0.93	0.83	0.61	0.89	0.62	0.68	0.33	0.78	0.73	0.84	0.82	-1	-0.33		-0.94	0.94	-0.08	0.91	-0.45
		0	0	0.01	0.04	0.2	0.02	0.19	0.14	0.53	0.07	0.1	0.01	0.04	0		0.37	0.23	0.21	0.95	0.27	0.7
Cyp1a2 RNAa	9.76	-0.82	-0.78			-0.57								-0.6	-0.63		0.68	0.72				-0.62
		0.01	0	0.06	0.02	0.24	0.24		0.07				0.01	0.21	0.56		0.53	0.49		0.33	0.11	0.58
S100A4 cells	8.89		0.85	0.64	0.96	0.86	0.55	0.31	0.92	0.43	0.32		0.25	-0.04	-0.97		-0.58					-0.22
		0.001	0	-	0.003			0.55		0.39	0.54	0.84	0.52	0.94		0.45			0.7	0.17		
Pparg RNAf	8.66	0.89	0.79	0.91		-0.18		0.87	0.47	0.4		-0.22		0.42		0.09			1		-0.22	7
		0.003	0	0.01		0.74			0.35	0.44	0.8	0.67	0.27	0.41	0.83		0.64		0.04	0.86	0.86	0.51
Cxcl5 RNAz	8.38	0.88	0.84	0.69		-0.45			0.55	0.3	0.5	0.37	0.69	0.47		0.22		-0.91			-0.15	
		0.004	0	0.13		0.37		0.28	0.26	0.56	0.32	0.47	0.04	0.35				0.27		0.9	0.91	0.9
Gsta2 RNAa			0.82		-0.07	0.71	0.84	0.75	0.71	0.29		-0.38		0.5	-0.89		0.28	1	0.71		-0.49	
C 0 00 DNA		0.0003	-	0.11	0.9			0.09	0.11	0.57	0.4	0.46	0.18	0.31		0.92		0.06		0.85	0.68	0.64
Cyp2c29 RNAa	8.17			-0.8		-0.15				-0.23	-0.4		-0.71			-0.42				-0.92		1
D. I. D. D. I.	0.00	0.002	0	0.05	0.11		0.19		0.84	0.67	0.43	0.16	0.03	0.69	0.99	0.72		0.26		0.26	0.99	0.03
Pde4a RNAf	8.09		0.59	0.6	0.31	-0.25	0.59	0.61	0.65	0.59	0.77	0.77	0.28	-0.25		-0.54		-0.68		0.7	0.74	-0.93
C II A DNIA	0.05		0.002	0.21	0.54			0.19	0.16	0.22	0.08	0.07	0.46	0.63	0.63	0.64	0.09		0.63	0.51	0.47	0.25
Cd14 RNAz	8.05	0.98	0.77	0.94	0.28	-0.16		-0.34	0.31	0.49	0.67	0.79	0.55	0.12	-0.56		1	-0.89	-0.1	0.77	0.98	-0.93
D 1 DNA£	7.0	0	0	0.01	0.59	0.76	0.42	0.51	0.55	0.33	0.14	0.06	0.12	0.82	0.62	0.65	0.04	0.3	0.94	0.44		0.24
Prom1 RNAf	7.9	0.91	0.63	0.34	-0.3	-0.5	0.7	0.6	0.87	0.66	0.65	0.64		0.05	-1	0.13		-0.98		0.94		-0.75
C 2-27 DNA-	7.05		0.001	0.51	0.56	0.31	0.12	0.21		0.15	0.17	0.17		0.92				0.14				0.46
Cyp2c37 RNAa	7.00	-0.92 0.001	-0.77		-0.92 0.01				-0.59 0.21	0.21	-0.58 0.23	-0.68 0.14		0.75	0.4	-0.44 0.71		-0.5 0.67	0.45	0.002		0.64
Informations	7.76	0.001	0.73	0.002	0.01			0.79		-0.27	0.23	0.14	0.11	0.75	0.4		0.72 -0.82		-0.99	0.08	0.06	0.56
Infarct area		0.90		0.0	0.74		0.45		0.39	0.6	0.37	0.07	0.09	0.30		0.21		0.30	0.07			0.83
II1b RNAz		0.89	0.77	-		-0.27			0.39	0.55	0.24	0.14		-0.22	n.a.	0.21	1	-0.77	0.07	0.95		-0.81
IIID KINAZ	1.14	0.003			0.43		0.22		0.74	0.33	0.33	0.49	0.47	0.68		0.76	0.05	0.44	0.72		0.11	0.4
Wisp1 RNAf	7.73		0.63			-0.47		0.64	0.74	0.63	0.76	0.49		0.02		-0.16		-1	0.49	0.55		-0.99
Wispi KitAi	1.10		0.001		0.01		0.11	0.17	0.02	0.00	0.78	0.06	0.40	0.02	0.73	0.10	0.46		0.62	0.0	0.43	
Gstm1 RNAa	7.7	0.86	0.77	-0.04		0.97	0.7	0.17		0.10		-0.55			-1	0.27	0.85	1		-0.79		0.29
Ostilii KivAa	1.1	0.00	0.77	0.95		0.001				0.00		0.33					0.35	0.04				
Timp1 RNAf	7.54	0.97	0.71	0.73	0.96		0.22			-0.17	0.61			-0.06		-0.02			-0.63			-0.99
impi ma	7.04	0.57	0.71		0.003		0.67	0.07		0.74	0.01	0.2	0.56	0.92			0.39	0.29	0.57			0.07
Ppara RNAa	7.51	-0.7	-0.57	-		-0.23			-0.75		-0.78			-0.2		-0.92					-0.95	
. para mini	7.01		0.003	0.1	0.18		0.32		0.09	0.2		0.26		0.71		0.25	0.53	0.41				0.37
Pdgfb RNAf	7.5	0.94	0.78	-		-0.67		0.28		0.24	0.66	0.67	0.45	0.06	-			-0.94				-0.76
6	0	0.0004		0.38	0.3	0.14				0.64	0.15			0.91				0.23				

4.12 CTGF cells

Factor	Cons		All											Progr				T18h		T2d	T5d	T14d
SMA cells	11.9	0.98	0.93	0.93	0.83	0.61	0.89	0.62	0.68	0.33	0.78	0.73	0.84	0.82	-1 0	-0.33 0.79	0.84	-0.94 0.23		0.08	0.91	-0.45 0.7
Cd14 RNAz	9.87		0.83	0.95	0.73	0.63	0.51	0.13	0.25			0.61	0.57	0.46	0.56	0.73	0.8		-0.42		0.27	0.08
		0	0	0.004	0.1	0.18	0.3	0.64	0.63	0.47	0.01	0.19	0.11	0.36	0.62	0.56	0.41	0.07		0.61	0.15	0.95
Ccr2 RNAz	9.64	0.95	0.89	0.84	0.7	0.23	0.44	-0.34	0.86	0.62	0.78	0.47	0.71	0.3	-0.27	1	0.78	0.97	-0.96	0.86	-0.65	-0.21
		0.0004	0	0.04	0.12	0.66	0.38	0.51	0.03	0.19	0.07	0.34	0.03	0.56	0.83	0.02	0.43	0.14	0.18	0.34	0.55	0.86
Cxcl5 RNAz	9.62		0.89	0.83	0.58	0.14	0.77	0.25	0.86	0.23	0.72		0.76	0.43	-0.65		0.98	1	0.12		-0.54	-0.95
		0.002	0	0.04	0.23	0.8	0.07	0.63	0.03	0.66	0.1	0.39		0.39	0.55	0.35	0.13	0.04		0.15	0.64	0.2
Ccl2 RNAz		0.96	0.92	0.84	0.77	0.43	0.11		0.58	0.19	0.78	0.59	0.82	0.62	-0.52		0.68		-0.99		-0.37	
C-IOC DNIA-		0.0002		0.03	0.08	0.39	0.83	0.37	0.23	0.72	0.07	0.22	0.01	0.19		0.64	0.52	0.11		0.44		0.86
Cd86 RNAz	9.40	0.93	0.85	0.47	0.38	0.3	0.85	0.51	0.91	0.7	0.75	0.41	0.6	-0.12 0.81	-0.75 0.46	0.99	0.77	0.03	0.79	0.75	-0.85 0.35	0.93
Timp1 RNAf	9.39		0.81	0.34	0.40	-0.14		-0.48	0.01	0.12	0.08	-	0.09	0.01	-0.7	0.11	0.38	-1	-0.85	1		0.55
Timpi Kivai	0.00	0.50	0.01	0.12	0.70	0.79	0.43	0.33	0.01	0.32		0.003		0.73	0.51	0.2	0.75	0.06		0.02	0.18	
Bili blood	9.28		0.91	0.53	0.83	0.67		-0.46		0.61	0.7	0.3	0.7	0.31	-0.24		-0.94					-0.93
	00	0.0001		0.28	0.04	0.15	0.72	0.36	0.11	0.2	0.12		0.04		0.85		0.21	0.46		0.93	0.55	0.25
II10rb RNAz	9.11	0.95	0.89	0.87	0.79	-0.01	0.33	-0.28	0.95	0.47	0.83	0.54	0.66	0.005	0.45	0.83	0.67	0.98	-0.87	0.51	-0.69	-0.62
		0.0004	0	0.02	0.06	0.99	0.53	0.59	0.004	0.35	0.04	0.27	0.06	0.99	0.71	0.38	0.54	0.12	0.33	0.66	0.52	0.57
Pdgfb RNAf	9.11	0.96	0.85	0.67	-0.01	-0.07	0.62	-0.04	0.97	0.51	0.9	0.95	0.59	0.25	0.84	1	0.96	1	-0.46	0.49	0.92	0.92
		0.0001		0.15		0.9	0.19		0.001			0.004		0.63		0.02		0.003				-
S100A4 cells	9.07	0.92	0.87	0.5	0.84	0.69	0.75	-0.44		0.8		-0.51		0.25		-0.87			-0.72			
T (I O DNIAC	0.04	0.001	0	0.31	0.04	0.13	0.09	0.39	0.02	0.05	0.09	0.3	0.89	0.64		0.33	0.98	0.05	0.49		0.18	
Tgfb2 RNAf	9.04	0.92	0.91	0.47	-0.58 0.22	-0.59 0.22	0.28	-0.25	0.95	-0.11 0.84	0.9	0.93	0.95 0.0001	0.91	0.94	0.9	0.05	0.06	-0.52 0.65		0.88	0.98
Tgfbr2 RNAf	8.93		0.9	0.55	-0.4	-0.61	0.59	-0.4	0.003	0.04	0.02	0.01	0.0001			-0.62	0.05		-0.79		0.98	0.14
Igibi Z KWAI		0.0002		0.26	0.43	0.01	0.85		0.0002				0.07		0.30		0.23	0.33	0.42		0.14	
II1b RNAz		0.92	0.85	0.86	0.81	0.21	0.5	0.55		-0.18			0.67		-0.69		0.88	0.95	0.45	0.75	-0.97	
		0.001	0	0.03	0.05	0.69	0.31	0.25	0.6	0.74				0.74		0.46	0.32		0.7	0.46	0.16	
Cyp2c37 RNAa	8.76	-0.9	-0.83	-0.9	-0.92	0.66	0.17	0.66	-0.65	-0.14	-0.87	-0.81	-0.69	-0.33	0.81	0.99	-0.13	0.77	0.71	1	-0.87	-0.97
• •		0.003	0	0.01	0.01	0.16	0.75	0.16	0.16	0.79	0.02	0.05	0.04	0.52	0.4	0.08	0.92	0.44	0.49	0.05	0.33	0.15
Actb RNAz	8.74	0.88	0.69	0.49	0.96	0.74	0.25	-0.48	0.82	0.66	0.78	0.84	0.07	-0.07	-0.59	0.99	-1	0.99	-1	0.97	0.84	0.33
		0.004			0.002	0.09	0.63	0.34	0.05	0.15	0.07	0.04	0.85	0.89	0.6		0.005	0.09	0.04		0.37	
Ifnar1 RNAz	8.7	0.94	0.87	0.61	0.87	0.73	0.14	-0.38	0.16	0.34	0.9	0.74	0.58	-0.32	-0.21	0.99	0.57	0.97	-0.97			-0.73
D DNAC		0.0004		0.2	0.02	0.1	0.78	0.46	0.76	0.52	0.01	0.09	0.1	0.53	0.87	0.11	0.61	0.14	0.15	0.58	0.57	0.48
Pparg RNAf	8.66		0.76	0.86	0.66	-0.54		0.47	0.8	-	-0.24	0.19	0.73	0.78	0.27			-0.56		0.99	0.2	0.95
		0.004	U	0.03	0.15	0.27	0.02	0.34	0.06	0.81	0.65	0.73	0.03	0.07	0.83	0.27	0.28	0.62	0.18	0.09	0.87	0.2

4.13 S100A4 cells

Factor	Cons	Αv	All	Init	6-18h	12-18h	12-30h	18-30h	Perp	30h-2d	B0h-5c	2-5d	2-14d	Progr	T0h	T6h	T12h	T18h	T30h	T2d	T5d	T14d
Gstm1 RNAa	9.38	0.93	0.89	0.3	0.77	0.8	0.94	0.89	0.87	0.75	0.29	-0.01	-0.08	0.11	0.98	0.83	-0.06	0.94	0.98	-0.6	-0.18	-1
		0.001	0	0.56	0.07	0.06	0.005	0.02	0.02	0.08	0.58	0.98	0.84	0.84	0.11	0.37	0.96	0.22	0.13	0.59	0.89	0.05
BrdU BE cells	9.3	0.82	0.86	0.82	0.35	-0.26	0.37	0.77	0.99	0.98	0.91	0.12	0.27	0.36	-0.49	1	0.88	0.11	0.54	1	0.25	0.75
		0.01	0	0.05	0.49	0.62	0.47			0.003		0.85	0.51	0.49	0.67	0	0.31	0.93	0.64	0	0.84	0.46
CTGF cells	9.07		0.87	0.5	0.84	0.69	0.75	-0.44		0.8		-0.51	0.05	0.25		-0.87	-0.04	-1	-0.72		-0.96	
Dill II	0.00	0.001	0	0.31	0.04	0.13	0.09	0.39	0.02	0.05	0.09	0.3	0.89	0.64	0.15		0.98	0.05	0.49	0.78	0.18	0.15
Bili blood	9.03		0.83	0.2	0.54	0.28	0.69	0.04	0.83	0.9	0.96	0.2	0.29	0.73			-0.29		0.66	0.97	0.84	-0.81
CMAII-	8.89	0.01	0.85	0.71	0.27	0.59	0.13	0.93	0.04	0.01	0.003		0.45	0.1	0.69	0.48		0.51	0.54	0.15	0.37	0.4
SMA cells	0.09	0.92	0.00	0.64	0.96	0.00	0.33	0.51	0.92	0.43	0.52	0.11		-0.04 0.94	-0.97 0.15		-0.58 0.61		-0.46 0.7	0.96	-0.99 0.09	
Cyp2c29 RNAa	8 25		-0.69	-	-0.83		-0.94			0.39	-0.7						-0.73			-0.78		
Cyp2C29 KWAa	0.23		0.0002	0.02	0.04	0.3	0.01	0.22	0.12	0.23	0.12	0.8	0.52	0.44		0.27	0.48	0.77	0.38	0.43	0.10	0.89
Gsta2 RNAa	8.23		0.8	-0.17	-0.1	0.5	0.76	0.63	0.9	0.77	0.78	0.17	0.09	0.17	0.75	0.75	0.63	• • • •	-0.95	0.03		-0.94
OSTAL THE TA	0.20	0.001	0.0	0.75	0.85	0.32	0.08	0.18	0.01	0.07	0.07		0.83	0.75	0.46	0.46	0.57	0.24	0.2	0.98	0.77	0.22
Cyp2c37 RNAa	8.13	-0.88	-0.76	-0.71	-0.96	0.16	-0.43		-0.78	-0.46	-0.76		0.09	0.34			-0.99		-1	-0.26	0.97	-0.89
-71		0.004	0	0.12	0.003	0.76	0.4	0.07	0.07	0.36	0.08	0.53	0.82	0.51	0.55	0.26	0.11	0.49	0.01	0.83	0.15	0.3
Cyp1a2 RNAa	8.09	-0.8	-0.77	-0.52	-0.9	-0.65	-0.91	-0.91	-0.59	0.14	-0.21	-0.08	-0.18	0.67	0.43	-0.12	0.21	0.49	-1	-0.71	1	-0.63
· .		0.02	0	0.29	0.01	0.16	0.01	0.01	0.22	0.79	0.69	0.87	0.64	0.14	0.72	0.92	0.87	0.67	0.05	0.5	0.02	0.56
Tnfrsf1a RNAz	7.9	0.83	0.78	-0.81	-0.38	0.05	0.79	-0.21	0.53	0.79	0.94	0.61	0.55	0.42	-0.17	-0.42	-0.94	-1	0.72	0.98	0.98	-0.73
		0.01	0	0.05	0.46	0.93	0.06	0.69	0.28	0.06	0.01	0.2	0.12	0.41	0.89	0.72	0.22	0.04	0.48	0.14	0.12	0.48
Bax RNAf	7.79	0.88	0.72	0.19		-0.48	0.6	0.61	0.89	0.77	0.62			-0.41	-0.72	-1	-0.97	-0.96	0.8	0.99	-0.28	0.58
		0.004	0	0.71	0.16	0.34	0.21	0.2	0.02	0.07	0.19	0.74	0.85	0.42	0.49	0.03	0.16	0.17	0.41	0.07		0.61
Tgfb1 RNAz	7.72		0.8			-0.48		0.8	0.92	0.81		-0.11		0.24			-0.62	-1	0.89	0.87		-0.43
D 111 ND 11	7.00	0.02	0	0.48	0.51	0.34	0.07	0.05	0.01	0.05	0.01	0.84	0.9	0.65	0.71	0.19	0.57	0.05	0.3	0.33		0.71
BrdU NP cells	7.62		0.79			-0.39		0.46	0.94	0.87	0.87	0.16	0.16	0.08		-0.71	1	-0.92			-0.38	
II10-l- DNA-	7 5 5	0.002	-	•	0.002		0.13	0.36	0.01	0.02	0.02	0.76	0.68	0.88	0.14	0.5	0.03	0.26	0.23	0.59	0.75	0.72
II10rb RNAz	7.55	0.78	0.79	0.11	0.35	-0.69 0.13	0.01	0.67	0.86	0.7	0.96	-0.12 0.82	0.02	0.14	0.65	-1 0.05	-0.77 0.44	-0.99 0.07	0.97	0.64	0.46	-0.79 0.42
Timp1 RNAf	7.41	0.02	0.72	0.005		-0.29		0.14	0.03	0.12	0.003		-0.38				-0.94				-0.84	
Tillipt KivAi	7.41	0.07	0.72	0.003	0.03	0.57	0.04	0.01	0.00	0.46	0.31	0.29	0.32	0.24	0.66	0.53	0.22	0.33	0.30	0.76	0.36	0.79
Col1a1 RNAf	7.38		0.74	-0.01	-0.7	-0.12		0.92	0.10	0.40				-	0.84	0.23	0.33	0.85	0.96		-0.66	
COILUI TATA	7.00	0.01	0.7 .	0.99	0.12		0.02	0.01	0.004		0.2	0.57	0.71	0.38	0.37	0.85	0.79	0.35	0.18	0.56	0.54	0.75
Slc10a1 RNAa	7.08	-0.88	_			-0.25					-0.77		-0.12	0.5	0.83	0.27	0.18		-0.36			-0.4
		0.004		0.71	0.36	0.63	0.27	0.64	0.15	0.04	0.08	0.71	0.76	0.31	0.38	0.83	0.89	0.09	0.77	0.15	0.15	0.74
Ccr2 RNAz	6.99	0.77	0.75	0.14	0.29	-0.51	0.77	0.68	0.51	0.29	0.94	-0.27	-0.02						0.89		0.41	-0.44
		0.03	0	0.79	0.57	0.3	0.07	0.14	0.3	0.58	0.01	0.6	0.95	0.51	0.98	0.36	0.55	0.19	0.3	0.44	0.73	0.71