

Plasma is preferred over serum for small RNA analysis using SEC

A) Schematic overview of methods used for plasma and serum isolation and subsequent experiments. B) Images of size-exclusion chromatography (SEC) before (left) and after (middle) applying plasma and flow-through (right). C) vtRNA1-1, miR127-3p, miR155-5p and miR451-5p in cHL patient plasma or serum after SEC. Data shown are from one representative donor (n=2) and depicted as mean Ct values +/- SEM of the 2 consecutive SEC fractions. D) Small RNA Bioanalyzer profiles of cHL (upper panel) and healthy donor (lower panel) plasma extracellular vesicles EV (left) and protein/HDL fraction (right). Profiles shown are of one representative donor (n=3).



RNAseq and particle analysis in plasma EVs and protein/HDL fractions

A) Distribution of small RNA sub classes in plasma protein/HDL fractions of a healthy individual (upper panel) and a cHL patient (lower panel). Data shown are of one representative donor (n=3) and depicted as % read counts. B) Distribution of individual miRNA species in protein/HDL fractions of a healthy donor (upper panel) and a cHL patient (lower panel). Data shown are of one representative donor (n=3) and depicted as % of miRNA read counts. C) Correlation between miRNA level and qNano particle concentration (left) and mode diameter (right) of healthy plasma extracellular vesicles (EVs) (upper panel) and cHL patient EVs (lower panel) (n=3 each). MiRNA level is based on the average Ct value (qRT-PCR) of miR127-3p, miR155-5p, miR21-5p and let7a-5p. r=Pearson correlation. D) RNAseg analysis of

miR142-3p, miR24-3p, miR155-5p and miR10b-5p in healthy and cHL patient plasma EVs (upper panel) and protein/HDL fractions (lower panel). Dots represent individual samples and lines are mean +/- SEM. Data are shown as summed read counts. †=not detected. E) Correlation between miRNA summed read counts and the number of different miRNAs detected by RNAseq in the plasma EV and protein/HDL fractions of healthy donors and cHL patients (n=12 samples). r=Pearson correlation. P value is calculated using an unpaired two tailed t test.



Candidate miRNA markers are disease specific

A-D) RT-PCR analysis of miR127-3p (A), miR155-5p (B), miR21-5p (C) and let7a-5p (D) in plasma extracellular vesicles (EVs) of healthy individuals (n=9), cHL patients (n=20) and SLE patients (n=4) after size-exclusion chromatography (SEC) and total RNA isolation using Trizol[®]. E-H) RT-PCR analysis of miR127-3p (E), miR155-5p (F), miR21-5p (G) and let7a-5p (H) in plasma EVs of male (n=6) and female (n=11) healthy individuals. For each individual the mean Ct value of SEC fractions 9 and 10 is used. Boxes show 25-75% percentile, whiskers show minimum-maximum and line represents the median. NS=not significant; * P<0.005; *** P<0.0005; **** P<0.0001 (unpaired two tailed t test).



Candidate miRNA levels in EVs from relapsed cHL and levels normalized to miR10b-5p

A-F) RT-PCR analysis of miR127-3p (A), miR155-5p (B), miR21-5p (C), let7a-5p (D), miR24-3p (E) and miR10b-5p (F) in plasma extracellular vesicles (EVs) of healthy individuals (n=9) and cHL patients with relapsed disease (n=7) after size-exclusion chromatography (SEC) and total RNA isolation using TRIzol[®]. G-K) RT-PCR analysis of miR127-3p (G), miR155-5p (H), miR21-5p (I), let7a-5p (J) and miR24-3p (K) in plasma EV of healthy individuals (n=7) and cHL patients (n=20), normalized to miR10b-5p. For each individual the mean Ct value of SEC fractions 9 and 10 is used. Boxes show 25-75% percentile, whiskers show minimum-maximum and line represents the median. NS=not significant; * P<0.05; ** P<0.005; *** P<0.0005 (unpaired two tailed t test).



MiR155-5p in EV outperforms total plasma and serum for distinguishing cHL patients from controls

A) RT-PCR analysis of miR155-5p in total plasma of healthy controls (n=7) and cHL patients (n=8) after RNA isolation using TRIzol-LS[®]. B) RT-PCR analysis of miR155-5p in extracellular vesicle (EV) fractions of the same healthy individuals and cHL patients as in (A) after size-exclusion chromatography (SEC) and total RNA isolation. For each individual the mean Ct value of SEC fractions 9 and 10 is used. C) RT=PCR analysis of miR155-5p in total serum of healthy individuals (n=8) and cHL patients (n=16). Boxes show 25-75% percentile, whiskers show minimum-maximum and line represents the median. NS=not significant; ** P<0.005 (unpaired two tailed t test).

Suppl Table 1

RNAseq data

%-age	58,36	10,35	3,09	2,67	2,33	1,85	1,63	1,60	1,15	49.82	13,63	3,85	3,26	2,72	2,39	2,22	2,16	1,92	1,34	005	3,93	3,85	1,57	1,43	1,31	1,31	1,24	1,04	25,85	13,66	11,94	6,83	5,00	3,99	ςΤ'ς 200	2,82	2,59	31,48	18,28	8,90 6 5 4	4.74	2,91	2,41	2,32	1,62	1,47	39,47	C/'7T	4.19	3.97	3,61	3,22	2,33	2,17 1,89
RC	106986	18981	5673	4887	4274	3397	2991	2925	2106	102868	28137	7954	6737	5611	4936	4590	4458	3968	2776	1162	914	895	366	333	304	304	288	241	140709	74359	64990	37157	27219	21708	1/143	15078	14081	279141	162100	78956	42022	25769	21390	20596	14365	13030	312111	C1001	33116	31413	28523	25431	18442	17135 14939
top 10 miRNA	hsa-miR-486-5p	hsa-miR-92a-3p	hsa-miR-16-5p	hsa-miR-26a-5p	hsa-miR-22-3p	hsa-miR-423-5p	hsa-miR-126-5p	hsa-let-7f-5p	hsa-miR-10b-5p	dc-TET-VIIII-BEII	hsa-miR-92a-3p	hsa-let-7a-5p	hsa-miR-26a-5p	hsa-let-7f-5p	hsa-miR-423-5p	hsa-miR-191-5p	hsa-miR-126-5p	hsa-miR-22-3p	hsa-mik-181a-5p hsa-mik-181a-bp	hterefordim-each	hsa-miR-191-5p	hsa-let-7a-5p	hsa-miR-26a-5p	hsa-miR-10b-5p	hsa-miR-423-5p	hsa-miR-126-5p	hsa-let-7f-5p	hsa-miR-22-3p	hsa-miR-486-5p	hsa-let-7a-5p	hsa-miR-92a-3p	hsa-let-7f-5p	hsa-miR-191-5p	hsa-miR-26a-5p	dc-524-XIM-BSN	hsa-mik-126-5p hsa-mik-22-3p	hsa-let-7b-5p	hsa-miR-486-5p	hsa-let-/a-bp	hsa-miR-191-5p hra-miP-00-2n	hsa-let-7f-5p	hsa-miR-26a-5p	hsa-miR-22-3p	hsa-miR-126-5p	hsa-let-7b-5p	hsa-miR-181a-5p	hsa-miR-486-5p	hra-mik-92a-5p	hsa-miR-26a-5p	hsa-let-7f-5p	hsa-miR-191-5p	hsa-miR-423-5p	hsa-let-7b-5p	hsa-miR-22-3p hsa-miR-126-5p
# different miRNAs	150									159	CCT								0	R									217									236									246							
miRNARC	183334									206465	0000								72751										544336									886790									790800							
total RC	521501									1018257	1770101								72766	0010									2611443									2350581									5452409							
sample	1									6	4)									1									2									n							
group	=	healthy donors	S							=	healthy donors	P							=	healthy donors	E								≥	cHL patients	R							2	CHL patients	EV							N otti mationte		2					
%-age	89,72	3,80	1,60	1,14	0,57	0,37	0,36	0,19	0,16	64.81	10,70	0,67	0,66	0,65	0,39	0,28	0,18	0,17	0,08	2 35	1,54	0,79	0,66	0,46	0,45	0,32	0,19	0,19	75,99	5,90	2,23	2,20	2,06	1,63	10/1	1,00	0,38	64,28	/4,11	3,59	1.86	1,25	1,20	1,17	66'0	0,91	91,44	1 25	0.86	0.72	0,57	0,47	0,20	0,18 0,15
ßC	980399	41493	17527	12479	6252	4052	3964	2086	1725	465055	4760	3274	3250	3198	1924	1358	869	848	408 670700	17673	11586	5939	4937	3498	3387	2386	1436	1416	91499	7102	2682	2645	2477	1964	9171	1206	460	17030	3066	951 710	492	331	319	309	262	242	115361	1571	1001	904	713	590	254	232 194
top 10 miRNA	hsa-miR-486-5p	hsa-miR-92a-3p	hsa-miR-16-5p	hsa-miR-423-5p	hsa-miR-22-3p	hsa-miR-25-3p	hsa-miR-10b-5p	hsa-miR-103a-3p	hsa-miR-126-5p	hca-miR-486-5n	hsa-miR-92a-3p	hsa-miR-423-5p	hsa-miR-191-5p	hsa-miR-10b-5p	hsa-miR-16-5p	hsa-miR-22-3p	hsa-miR-10a-5p	hsa-miR-25-3p	hsa-miR-181a-5p hsa-mip-46-5p	de-eco-dim-esq	hsa-miR-16-5p	hsa-miR-423-5p	hsa-miR-22-3p	hsa-miR-191-5p	hsa-miR-10b-5p	hsa-miR-25-3p	hsa-miR-103a-3p	hsa-miR-10a-5p	hsa-miR-486-5p	hsa-miR-92a-3p	hsa-miR-22-3p	hsa-miR-191-5p	hsa-miR-423-5p	hsa-miR-16-5p	dc-q7T-XIM-BSN	hsa-miR-26a-50 hsa-miR-10b-50	hsa-miR-146a-5p	hsa-miR-486-5p	dd-191-Aim-Ban	hsa-miR-92a-3p hra-miP-22-3p	hsa-miR-26a-5p	hsa-miR-423-5p	hsa-miR-10b-5p	hsa-miR-126-5p	hsa-let-7a-5p	hsa-miR-10a-5p	hsa-miR-486-5p	dc-676-Alm-esh	hsa-miR-191-5p	hsa-miR-16-5p	hsa-miR-10b-5p	hsa-miR-22-3p	hsa-miR-25-3p	hsa-miR-126-5p hsa-miR-26a-5p
# different miRNAs	109									75	2								114	-									102									67									61							
miRNARC	1092693									490501	Toront								757704										120404									26493									126160							
totalRC	1318571									594752									047257										312559									83318									190459							
sample	1									6	4)									1									2									n							
group	_	healthy donors	protein/HDL							_	healthy donors	protein/HDL							-	healthy donors	protein/HDL								=	cHL patients	protein/HDL							=	CHL patients	protein/HDL							II otti mationte	urctein/HDI						