

Supplementary Information

Supplementary Tables

Table S1. CCC and overall CCC with bootstrap 95% CI (all human miRNAs).

<i>Sample</i>	<i>Pair</i>	<i>CCC</i>	<i>Affymetrix</i>		<i>CCC</i>	<i>Agilent</i>		<i>CCC</i>	<i>Illumina</i>	
			<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>
<i>hREF</i>	1–2	0.988			0.997			0.991		
	1–3	0.993	0.992	(0.990–0.993)	0.989	0.994	(0.993–0.995)	0.993	0.994	(0.994–0.995)
	2–3	0.994			0.995			0.999		
<i>A498</i>	1–2	0.939			0.975			0.996		
	1–3	0.893	0.93	(0.911–0.943)	0.970	0.975	(0.969–0.979)	0.981	0.989	(0.987–0.991)
	2–3	0.961			0.962			0.989		

Table S2. CCC and overall CCC with bootstrap 95% CI (all miRNAs).

<i>Sample</i>	<i>Pair</i>	<i>CCC</i>	<i>Affymetrix</i>		<i>CCC</i>	<i>Agilent</i>		<i>CCC</i>	<i>Illumina</i>	
			<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>
<i>hREF</i>	1–2	0.990			0.997			0.990		
	1–3	0.995	0.993	(0.993–0.994)	0.990	0.994	(0.994–0.995)	0.992	0.994	(0.993–0.994)
	2–3	0.996			0.996			0.999		
<i>A498</i>	1–2	0.959			0.977			0.996		
	1–3	0.928	0.954	(0.950–0.957)	0.970	0.976	(0.970–0.980)	0.981	0.988	(0.987–0.990)
	2–3	0.976			0.980			0.988		

Table S3. CCC and overall CCC with bootstrap 95% CI (quantile normalization).

<i>Sample</i>	<i>Pair</i>	<i>CCC</i>	<i>Affymetrix</i>		<i>CCC</i>	<i>Agilent</i>		<i>CCC</i>	<i>Illumina</i>	
			<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>		<i>OCCC</i>	<i>CI 95%</i>
<i>hREF</i>	1–2	0.992			0.998			0.999		
	1–3	0.994	0.994	(0.992–0.995)	0.998	0.998	(0.997–0.998)	0.998	0.999	(0.998–0.999)
	2–3	0.995			0.998			0.999		
<i>A498</i>	1–2	0.972			0.991			0.998		
	1–3	0.967	0.971	(0.963–0.977)	0.993	0.993	(0.991–0.994)	0.995	0.996	(0.995–0.997)
	2–3	0.973			0.994			0.995		

Table S4. CCC and overall CCC with bootstrap 95% CI (loess normalization).

<i>Sample</i>	<i>Pair</i>	<i>Affymetrix</i>			<i>Agilent</i>			<i>Illumina</i>		
		<i>CCC</i>	<i>OCCC</i>	<i>CI 95%</i>	<i>CCC</i>	<i>OCCC</i>	<i>CI 95%</i>	<i>CCC</i>	<i>OCCC</i>	<i>CI 95%</i>
<i>hREF</i>	1–2	0.989			0.998			0.998		
	1–3	0.991	0.992	(0.989–0.993)	0.997	0.998	(0.997–0.998)	0.998	0.998	(0.998–0.999)
	2–3	0.994			0.998			0.999		
<i>A498</i>	1–2	0.983			0.990			0.998		
	1–3	0.978	0.981	(0.976–0.985)	0.992	0.992	(0.990–0.994)	0.995	0.996	(0.995–0.997)
	2–3	0.981			0.994			0.995		

Table S5. Estimates of the linear measurement error model, $\lambda = 1$ (quantile normalization).

<i>Sample</i>	<i>Pair</i>	<i>Estimate</i>	a_0		b_0	
			<i>Estimate</i>	<i>CI 95%</i>	<i>Estimate</i>	<i>CI 95%</i>
<i>hREF</i>	Agilent vs. Affymetrix	–8.1304	(–8.4960, –7.7648)	2.2040	(2.1378, 2.2702)	
	Illumina vs. Affymetrix	–8.3585	(–9.0694, –7.6476)	2.8994	(2.8071, 2.9918)	
	Illumina vs. Agilent	2.9633	(2.7599, 3.1668)	1.1761	(1.1204, 1.2319)	
<i>A498</i>	Agilent vs. Affymetrix	–9.0705	(–9.5760, –8.5650)	2.3613	(2.2836, 2.4391)	
	Illumina vs. Affymetrix	–11.2364	(–12.5018, –9.9710)	3.4002	(3.2772, 3.5233)	
	Illumina vs. Agilent	2.8587	(2.3850, 3.3325)	1.2101	(1.1250, 1.2952)	

Table S6. Estimates of λ and CI 95% (quantile normalization). Values obtained as a ratio of σ_ϵ^2 (error variance of Y) and σ_δ^2 (error variance of X), estimated via random effects models.

<i>Sample</i>	<i>Pair</i>	λ	<i>CI 95%</i>
<i>hREF</i>	<i>Agilent-Affymetrix</i>	3.116	2.878–3.374
	<i>Illumina-Affymetrix</i>	3.891	3.594–4.213
	<i>Illumina Agilent</i>	1.249	1.153–1.352
<i>A498</i>	<i>Agilent-Affymetrix</i>	3.132	2.892–3.390
	<i>Illumina-Affymetrix</i>	3.827	3.535–4.144
	<i>Illumina-Agilent</i>	1.222	1.129–1.323

Table S7. Estimates of the linear measurement error model, λ estimated (quantile normalization).

<i>Sample</i>	<i>Pair</i>	a_0		b_0	
		<i>Estimate</i>	<i>CI 95%</i>	<i>Estimate</i>	<i>CI 95%</i>
<i>hREF</i>	Agilent vs. Affymetrix	-5.6388	(-5.7125, -5.5650)	1.7689	(1.7392, 1.7987)
	Illumina vs. Affymetrix	-3.0838	(-3.1592, -3.0085)	1.9784	(1.9483, 2.0084)
	Illumina vs. Agilent	3.2250	(3.0627, 3.3873)	1.1179	(1.0680, 1.1677)
<i>A498</i>	Agilent vs. Affymetrix	-5.8006	(-5.8875, -5.7138)	1.7923	(1.7600, 1.8245)
	Illumina vs. Affymetrix	-3.1551	(-3.2488, -3.0614)	1.9938	(1.9604, 2.0273)
	Illumina vs. Agilent	3.3203	(2.9651, 3.6755)	1.1075	(1.0338, 1.1812)

Table S8. Number (n) and proportion (%) of miRNA lying within the agreement intervals, estimated according to the measurement error model parameters estimated by setting $\lambda = 1$ and by estimating it via random effects models. Data were normalized according to the quantile normalization algorithm. Confidence intervals for the proportions were computed using the Clopper–Pearson exact method.

<i>Sample</i>	<i>Comparison</i>	$\lambda = 1$		λ Estimated	
		% (<i>CI</i> 95%)	<i>n</i>	% (<i>CI</i> 95%)	<i>n</i>
<i>hREF</i>	<i>Agilent-Affymetrix</i>	79.46 (76.52, 82.19)	646	81.55 (78.71, 84.16)	663
	<i>Illumina-Affymetrix</i>	71.34 (68.1, 74.43)	580	80.93 (78.06, 83.58)	658
	<i>Illumina-Agilent</i>	96.43 (94.92, 97.6)	784	96.19 (94.63, 97.39)	782
<i>A498</i>	<i>Agilent-Affymetrix</i>	78.84 (75.87, 81.6)	641	82.41 (79.62, 84.97)	670
	<i>Illumina-Affymetrix</i>	70.48 (67.21, 73.6)	573	83.89 (81.18, 86.35)	682
	<i>Illumina-Agilent</i>	98.52 (97.44, 99.24) †	801	98.77 (97.75, 99.41) †	803

†: the platform pair is in agreement.

Table S9. Estimates of the linear measurement error model, $\lambda = 1$ (loess normalization).

<i>Sample</i>	<i>Pair</i>	a_0		b_0	
		<i>Estimate</i>	<i>CI 95%</i>	<i>Estimate</i>	<i>CI 95%</i>
<i>hREF</i>	Agilent vs. Affymetrix	-8.5056	(-8.8715, -8.1397)	2.2852	(2.2189, 2.3515)
	Illumina vs. Affymetrix	-8.7511	(-9.4867, -8.0155)	2.9867	(2.8927, 3.0807)
	Illumina vs. Agilent	3.0661	(2.8601, 3.2720)	1.1540	(1.0984, 1.2096)
<i>A498</i>	Agilent vs. Affymetrix	-8.6769	(-9.1779, -8.1759)	2.2718	(2.1943, 2.3493)
	Illumina vs. Affymetrix	-11.0793	(-12.3307, -9.8279)	3.3492	(3.2267, 3.4718)
	Illumina vs. Agilent	2.6593	(2.1942, 3.1245)	1.2559	(1.1700, 1.3418)

Table S10. Estimates of λ and CI 95% (loess normalization). Values obtained as ratio of σ_ϵ^2 (error variance of Y) and σ_δ^2 (error variance of X), estimated via random effects models.

<i>Sample</i>	<i>Pair</i>	λ	CI 95%
hREF	<i>Agilent-Affymetrix</i>	3.306	3.054–3.579
	<i>Illumina-Affymetrix</i>	4.021	3.714–4.354
	<i>Illumina Agilent</i>	1.216	1.123–1.317
A498	<i>Agilent-Affymetrix</i>	2.967	2.740–3.212
	<i>Illumina-Affymetrix</i>	3.779	3.490–4.091
	<i>Illumina-Agilent</i>	1.274	1.176–1.379

Table S11. Estimates of the linear measurement error model, λ estimated (loess normalization).

<i>Sample</i>	<i>Pair</i>	a_0		b_0	
		<i>Estimate</i>	CI 95%	<i>Estimate</i>	CI 95%
hREF	<i>Agilent vs. Affymetrix</i>	−5.8600	(−5.9285, −5.7916)	1.8232	(1.7945, 1.8518)
	<i>Illumina vs. Affymetrix</i>	−3.1706	(−3.2438, −3.0973)	2.0121	(1.9825, 2.0418)
	<i>Illumina vs. Agilent</i>	3.2998	(3.1301, 3.4694)	1.1030	(1.0525, 1.1534)
A498	<i>Agilent vs. Affymetrix</i>	−5.6135	(−5.7061, −5.5208)	1.7369	(1.7036, 1.7703)
	<i>Illumina vs. Affymetrix</i>	−3.1723	(−3.2666, −3.0780)	1.9687	(1.9350, 2.0023)
	<i>Illumina vs. Agilent</i>	3.2046	(2.8815, 3.5276)	1.1301	(1.0585, 1.2017)

Table S12. Number (n) and proportion (%) lying within the agreement intervals, estimated according to the measurement error model parameters by setting $\lambda = 1$ and via random effects models. Data were normalized according to the loess normalization algorithm. Confidence intervals for the proportions were computed using the Clopper–Pearson exact method.

<i>Sample</i>	<i>Comparison</i>	$\lambda = 1$		λ Estimated	
		% (CI95%)	n	% (CI95%)	n
hREF	<i>Agilent-Affymetrix</i>	79.21 (76.26, 81.95)	644	81.18 (78.32, 83.81)	660
	<i>Illumina-Affymetrix</i>	71.83 (68.60, 74.90)	584	82.90 (80.14, 85.43)	674
	<i>Illumina-Agilent</i>	96.06 (94.49, 97.29)	781	95.82 (94.2, 97.09)	779
A498	<i>Agilent-Affymetrix</i>	79.95 (77.03, 82.65)	650	83.27 (80.53, 85.77)	677
	<i>Illumina-Affymetrix</i>	68.76 (65.45, 71.93)	559	83.03 (80.27, 85.54)	675
	<i>Illumina-Agilent</i>	98.4 (97.28, 99.15)†	800	98.52 (97.44, 99.24)†	801

† the platform pair is in agreement.

Supplementary Figures

Figure S1. Box and density plots for both samples. The left column refers to hREF and the right column to A498. Plots refer to quantile normalized log₂-transformed data. (**Lower panels**) The solid line represents the technical replicate labeled as 1 in the datasets, whereas the dashed line and dotted line represent technical Replicates 2 and 3, respectively.

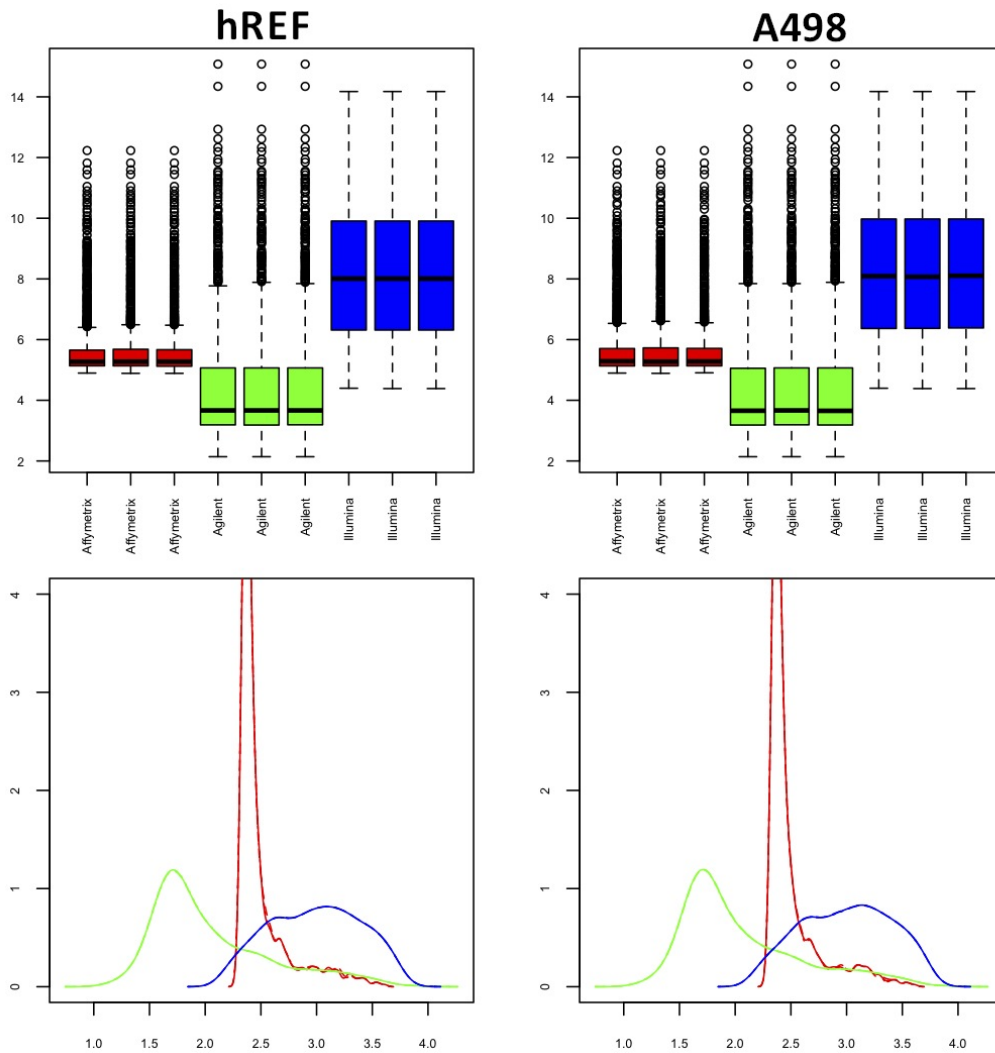


Figure S2. Box and density plots for both samples. The left column refers to hREF and the right column to A498. Plots refer to loess normalized log₂-transformed data. (**Lower panels**) The solid line represents the technical replicate labeled as 1 in the datasets, whereas the dashed line and dotted line represent technical Replicates 2 and 3, respectively.

