

Millipore

analyte	T0serum_PCT	T0heparin_PCT	T3edta_PCT	T3heparin_PCT	T0serum_median	T0heparin_median
AMYLIN_I	29	27	40	34	47.01	36.95
BCA_1	100	100	100	100	22.42	29.135
CCL19_M	100	100	100	100	117.685	143.03
CCL20_M	67	89	80	78	5.78	7.12
CKINE	69	62	85	78	428.95	436.75
CRP	100	100	100	100	5484776.395	6192559.445
CTACK	100	100	100	100	637.905	550.33
CXCL11_I	99	100	98	100	41.68	374.545
CXCL6_G	100	100	100	100	104.14	123.28
CXCL9_M	100	100	100	100	1803.465	1727.775
C_PEPTIC	99	99	99	100	1940.46	1999.9
EGF	93	56	49	60	96.45	20.02
ENA_78	100	99	99	100	1096.2	351.91
EOTAXIN	100	100	100	100	108.105	413.075
EOTAXIN_	100	100	100	100	961.485	378.715
EOTAXIN_	26	96	14	92	15.375	36.22
FGF_BAS	99	99	99	100	504.34	160.27
FIT_3_LIG	61	56	58	54	35.02	34.115
FRACTAL	32	32	53	30	43.43	58.2
GHRELIN	28	9	17	26	30.005	20.91
GIP	99	100	99	96	33.73	71.045
GLP_1	50	42	72	54	44.3	54.98
GLUCAG(C	41	49	56	58	41.57	36.19
GM_CSF	97	98	98	98	22.18	19.095
GRO	100	100	100	100	1068.99	820.225
G_CSF	65	83	81	82	25.02	38.91
IFNA2	41	37	56	46	14.49	18.98
IFNG	79	81	87	82	10.5	9.28
IL_10	99	94	88	96	8.5	3.545
IL_11	83	100	76	100	89.06	180.06
IL_12P40	69	68	70	62	58.25	74.57
IL_12P70	44	52	74	42	3.68	4.12
IL_13	31	27	36	34	3.76	2.96
IL_15	72	80	90	88	2.825	3.285
IL_16	73	74	89	84	67.15	72.36
IL_17	31	52	81	56	3.36	3.48

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IL_1A	96	100	99	100	35.485	70.125
IL_1B	67	67	66	46	3.68	3.37
IL_1RA	67	74	85	72	12.37	11.865
IL_2	56	49	55	48	1.97	1.77
IL_20	23	18	42	20	187.14	150.315
IL_21	3	3	2	2	13.67	7.67
IL_23	55	52	56	58	211.59	191.27
IL_28A	14	12	50	20	36.815	22.32
IL_29_IFN	50	66	46	64	166.34	182.96
IL_3	9	15	15	28	4.21	9.16
IL_33	39	39	80	48	32.73	32.53
IL_4	1	5	1	20	531.48	18.09
IL_5	12	14	16	14	1	0.865
IL_6_1	86	83	90	86	5.225	5.11
IL_7	58	41	44	42	6.745	5.92
IL_8	100	99	100	100	15.025	10.2
IL_9	31	31	31	30	4.84	4.21
INSULIN	97	97	99	98	675.07	686.67
IP_10	100	100	100	100	396.975	698.23
I_309	80	78	91	86	2.72	2.48
LEPTIN	100	100	100	100	13728.36	12629.69
LIF	55	58	80	66	13.2	10.205
MCP_1	100	100	100	100	547.56	404.345
MCP_2	99	92	100	96	51.09	22.355
MCP_3	75	65	76	76	17.85	16.96
MCP_4	60	42	47	50	102.955	107.29
MDC	100	100	100	100	1235.43	687.185
MIP_1A	64	65	67	62	31.345	28.44
MIP_1B	100	99	99	98	51.18	38.1
MIP_1D	100	100	99	98	2711.49	2843.335
M_CSF	2	0	0	0	1744.845	
PP	95	96	97	98	120.52	172.575
PYY	91	87	97	92	102.77	98.74
SAA	100	99	100	100	7531249.87	7080872.77
SAP	100	100	100	100	45844262.11	46194714.94
SCD30	38	34	38	40	40.39	54.19
SCD40L	46	98	95	92	50414.42	2534.485

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SCF	79	85	87	90	30.38	30.86
SDF_1A	98	98	99	100	2113.185	1727.6
SEGFR	100	100	100	100	44621.015	48619.91
SGP130	100	100	100	100	202861.015	183221.81
SILRII	100	100	100	100	6934.79	7078.945
SIL_1RI	48	57	48	52	45.685	45.57
SIL_2RA	65	62	62	64	46.26	49.635
SIL_4R	82	83	74	76	375.8	355.71
SIL_6R	100	100	100	100	20603.27	21502.305
SRAGE	37	54	33	56	34.67	38.815
STNFRI	100	99	100	100	1207.88	1083.1
STNFRII	100	100	100	100	5738.595	5669.22
SVEGFR1	54	61	67	54	795.575	1034.17
SVEGFR2	100	100	100	100	7408.92	8173.165
SVEGFR3	47	49	57	62	1571.5	1437.09
TARC	100	99	99	98	125.235	24.63
TGFA	87	56	50	48	6.02	3.755
TNFA_1	99	99	98	98	8.23	7.73
TNFB	29	41	70	34	8.7	6.98
TPO	97	96	93	96	304.04	187.405
TRAIL	76	83	75	74	33.155	39.38
TSLP	26	30	33	36	11.52	10.09
VEGF	95	53	48	36	162.52	51.18
XCL1_LY†	17	31	24	28	63.33	53.33

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T3edta_median	T3heparin_median	T0serum_within_CV	T0heparin_within_CV	T3edta_within_CV	T3heparin_within_CV
49.89	45.51	1.4906	2.5835	2.4526	1.2447
25.3	34.29	11.7991	10.0446	3.1081	2.0949
119.16	132	3.1869	3.2187	1.6943	1.9649
6.425	6.32	6.8112	5.8892	6.1992	8.1761
439.86	358.69	5.7796	5.9968	9.1422	6.4999
7420717.195	8292279.14	1.2753	1.1268	1.3104	2.9279
620.69	550.785	1.8423	1.4226	0.7949	0.9754
26.925	404.62	22.5867	12.6593	2.9988	1.3445
45.815	93.18	2.635	3.5075	2.5127	3.0491
1819.28	2284.15	5.1018	4.9969	1.6353	1.406
2512.345	2453.04	1.2936	2.3139	0.7725	0.8179
28.01	28.735	2.0289	12.2421	17.0151	17.6586
359.58	352.61	5.448	5.4993	2.4121	1.444
50.16	407.03	10.6487	4.8203	2.0381	1.4701
388.87	339.345	4.5672	4.5462	1.3516	1.7863
22.785	36.255	2.6776	3.9305	14.9746	2.0197
116.98	127.605	5.4308	7.6044	4.161	2.6692
24.45	32.67	11.8752	13.001	9.2515	21.6828
78.33	121.02	18.4918	12.1174	17.29	0.6402
32.155	32.79	7.7809		4.3993	
118.32	168.06	9.1896	6.078	2.3443	1.4879
51.2	90.38	3.5367	15.694	4.6521	11.5488
55.52	42.16	2.4705	3.6183	3.2288	4.0081
17.55	20.12	17.3595	9.2024	9.7291	5.7042
854.58	849.09	1.241	1.2678	2.417	1.205
36.96	49.76	24.6852	6.3389	10.6013	6.5778
16.91	20.59	23.0741	43.1071	26.2101	8.8511
11.07	9.98	30.0694	16.7799	14.2318	14.0207
3.575	2.605	15.5586	24.5329	18.4839	31.6568
48.09	182.005	10.2316	5.0302	5.5584	2.8625
62.9	91.65	4.5804	7.0536	7.3899	4.8462
4.3	5.35	22.5499	54.2486	35.5848	42.6684
3.09	3.09	75.8955	6.4198	46.1792	
3.325	3.45	17.2226	25.527	24.6025	24.5557
82.71	76.675	14.5537	8.0709	7.1665	6.4858
5.47	2.62	45.1008	51.786	24.3317	36.4594

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62.565	85.58	19.8394	10.9979	4.986	6.883
2.86	5.62	34.2455	34.8569	24.7859	23.725
14.25	10.66	10.8018	26.3265	9.4975	11.403
1.7	2.15	67.0333	45.0481	32.1829	46.4301
277.99	458.81	1.2158	19.092	5.0677	0
	7.97				
148.1	200.12	16.714	9.242	3.9168	4.2643
18.49	70.175	7.4344	7.7108	9.4737	3.5183
139.46	199.605	8.1762	4.2264	4.932	8.1924
8.47	12.035	19.6327	19.455	19.0706	2.7285
26.54	27.76	5.1313	15.4623	8.798	12.3058
616.57	41.07			0.1493	18.9537
1.26	1.62	19.3031	0	94.9071	
4.39	4.7	37.6052	33.5626	16.5249	22.6234
6.51	8.35	26.3939	17.6007	17.201	14.3214
5.895	8.255	5.797	10.7446	10.1123	5.954
4.69	5.36	9.9944	38.9868	24.5058	35.4517
995.84	1226.79	4.45	6.2707	1.0612	0.9865
570.685	761.68	6.6967	6.0166	1.6938	2.3304
2.77	2.88	35.9288	26.6865	16.1186	24.3977
10685.345	12083.04	1.0321	2.8878	0.5302	0.7298
14.46	11.9	11.8027	11.9199	9.9488	6.0407
253.48	406.505	0.8999	1.1179	1.7053	0.6535
28.06	24.71	5.9144	7.9877	6.8702	9.4812
14.57	16.98	21.1619	12.9772	19.6698	9.0422
84.34	88.06	6.74	7.3866	5.8138	3.0403
876.475	665.56	3.046	3.152	1.7953	3.4598
25.09	28.01	3.8237	9.5192	13.5651	6.3947
33.84	36.57	5.8252	6.5628	6.7006	2.9243
2491.25	3441.74	2.1505	2.2429	1.2796	1.4134
172.21	209	2.3599	3.6004	2.1404	1.1274
85.65	98.405	7.2467	6.2474	2.5216	1.4507
8510039.73	10378829.09	1.5034	1.1607	0.9209	1.5138
43113636.28	46379519.59	1.038	0.6718	0.7914	0.9685
57.635	46.88	7.7698	4.7421	7.3629	1.6331
993.865	1590.535	15.2958	12.2171	6.3119	2.431

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27.97	30.74	10.3704	4.1851	10.8106	6.2174
1999.28	1987.32	3.6196	1.7332	1.4929	1.4405
41520.765	47039.885	0.8787	0.9175	1.3195	1.0469
165099.42	184946.205	0.9809	1.7183	2.4644	1.3821
5833.77	6177.6	1.0136	0.7704	1.0909	0.6139
45.12	50.31	3.0522	2.3956	5.4572	5.966
51.78	38.19	5.1962	15.5133	27.6895	10.953
391.12	345.215	3.7862	5.1637	4.2856	6.8601
18589.51	21087.26	0.9198	0.6147	0.7026	0.4818
36.57	38.185	3.6057	4.5581	7.3576	4.3202
960.92	1179.8	1.4783	1.6153	1.5436	0.9209
5020.485	5799.12	1.259	1.2846	1.717	1.3921
823.265	767.36	8.7661	9.8547	6.8061	7.2606
6236.505	7535.17	1.8815	2.587	2.5542	2.7227
1587.3	2395.73	13.9654	12.0561	5.2547	11.007
27.44	26.35	9.5562	13.0285	3.4322	7.755
3.565	3.7	9.3755	35.2758	23.3526	36.9527
9.135	8.1	12.472	20.2159	14.4442	11.2669
11.85	6.47	12.9089	23.2118	16.2926	44.3366
122.98	169.405	8.0748	5.2698	4.9739	6.2362
23.265	43.28	7.855	9.0511	17.652	11.7947
5.53	7.795	10.7897	24.1442	13.1105	8.5944
53.37	82.58	2.1657	17.4602	7.8764	9.6107
41.77	52.005	9.3641	8.3279	11.3371	8.6244

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T0serum_across_CV	T0heparin_across_CV	T3edta_across_CV	T3heparin_across_CV	T0serum_within_ICC
10.8309	8.7455	8.9612	7.2588	0.98
10.4637	7.0579	5.9516	3.8565	0.87
6.4942	5.433	4.3892	2.8996	0.97
6.6444	11.0193	12.0602	10.3508	0.97
10.8503	7.4256	11.2702	6.7479	0.83
1.634	1.2014	1.8649	0.6197	0.99
4.2455	4.634	2.8989	2.2727	0.96
7.0034	10.3354	6.5928	2.8461	0.8
4.7137	3.3642	6.2814	4.7827	0.94
4.9182	2.6903	2.3898	2.937	0.93
2.7043	3.563	1.772	1.3881	0.99
8.2921	10.786	17.2579	4.2034	1
2.977	4.1567	2.4841	1.5359	0.87
4.6843	7.3109	8.5416	8.6107	0.7
4.5642	5.2522	3.2612	2.8222	0.9
0.6676	7.0569	7.1608	3.8347	0.98
3.2228	4.3475	7.1235	2.7465	0.71
9.5271	12.429	23.7268	7.5558	0.92
50.0373	11.8628	11.9496	2.2236	0.85
6.3959	3.5379	0.319	22.5865	0.75
5.867	6.5373	2.3213	5.2514	0.93
6.6869	7.6504	8.3419	11.5679	0.99
12.7747	29.4693	14.0922	9.8502	0.98
13.7654	26.4688	21.2285	15.5527	0.82
2.2376	3.0955	2.5592	2.0282	0.97
11.8756	10.4462	14.749	7.4483	0.69
24.8953	15.8634	16.0219	12.8628	0.85
16.0229	25.185	15.2234	18.8354	0.83
15.5779	21.0152	22.7376	38.0758	0.89
12.5426	10.5846	14.9559	3.1037	0.84
8.5467	8.2999	6.8054	6.3306	0.99
35.6438	99.704	44.2341	39.8108	0.92
25.3888	23.8164	38.4309	19.6773	0.55
63.4967	53.8989	37.1805	11.5843	0.98
13.9631	16.8036	9.4572	10.8018	0.72
31.9477	58.5247	23.897	31.8065	0.88

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18.4423	12.3847	10.1244	7.4531	0.74
63.4438	65.7644	46.8909	17.9801	0.93
39.3731	24.1518	25.9986	22.5995	0.98
50.4189	71.4043	62.1136	1.7311	0.82
15.0346	12.6979	6.1982	12.3233	1
	8.386			
20.3253	25.8729	12.1168	14.6948	0.86
		14.5504	9.1492	0.96
14.0307	10.128	4.1654	4.5886	0.84
	25.1585	6.0547	42.1476	0.92
11.7532	22.7271	9.4242	12.0788	0.99
			2.9824	
-9.8036	388.4029	132.4	0.5572	0.98
40.8214	24.1374	29.883	18.4643	0.81
46.1724	26.1355	18.4129	6.1979	0.82
10.0294	15.2317	12.7598	11.6193	0.98
56.6226	50.3424	34.2205	7.0692	0.98
3.3296	10.2738	3.6459	2.0746	0.95
3.7032	3.3232	2.4034	6.0118	0.81
70.7723	43.466	30.5502	40.7628	0.85
2.1244	2.0204	1.5869	1.0339	0.99
18.3793	11.684	14.3089	16.7454	0.81
2.2651	1.9048	2.8984	1.7343	0.99
7.3257	17.1592	11.4753	8.896	0.92
17.6879	35.7903	14.3609	23.106	0.82
6.8598	7.9237	7.7976	18.3784	0.9
2.2686	3.882	2.521	2.5543	0.82
7.9567	11.1593	19.0922	6.5078	0.99
5.4191	11.6214	5.0648	3.7677	0.91
5.4268	5.0333	6.7994	4.1287	0.97
3.4156	3.799	2.3156	1.7902	0.98
4.2149	5.3339	4.9875	2.4905	0.81
2.9939	1.2418	1.8457	1.3165	0.97
1.6237	0.832	1.6926	0.7542	0.84
11.1771	37.7924	14.561	4.6697	0.9
5.4623	4.0625	8.121	5.0518	0.53



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15.6415	8.9144	10.9816	6.2532	0.9
4.5731	3.9737	5.1151	2.0698	0.83
3.9867	7.4142	6.3794	1.0464	0.97
4.2082	6.0598	6.6932	3.6281	0.96
4.5816	5.5906	8.1543	1.3712	0.94
5.4113	7.7728	7.2444	4.2755	0.96
12.0641	9.7605	17.0067	3.1033	0.97
10.4246	11.6592	13.9141	4.5638	0.89
4.0475	6.5083	7.0507	0.7338	0.98
1.6868	5.3447	4.5331	15.6118	0.97
6.4424	7.6911	9.9077	0.9852	0.98
5.4291	7.2065	8.9889	2.216	0.98
7.8943	11.9469	14.6119	11.1877	0.74
5.6504	6.0233	8.9433	1.7629	0.9
9.2985	10.9004	12.1915	3.8815	0.46
5.3428	13.56	7.1192	5.0035	0.88
24.2494	54.7498	29.9678	5.1322	0.96
16.0385	24.5439	14.1867	15.1972	0.91
77.1184	33.3719	13.3148	8.9386	0.92
6.8721	13.3921	7.9788	6.1192	0.86
16.6658	16.703	16.6314	12.8186	0.96
15.4176	36.1937	11.6511	5.5913	0.93
5.4496	8.5454	19.8113	2.5934	0.98
	14.0632		6.9169	0.38

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T0heparin_within_ICC	T3edta_within_ICC	T3heparin_within_ICC	T0serum_across_ICC	T0heparin_across_ICC
0.82	0.97	0.99	0.62	0.53
0.77	0.98	0.99	0.84	0.84
0.95	0.99	0.99	0.82	0.93
0.98	0.98	0.98	0.99	0.95
0.94	0.82	0.83	0.7	0.82
0.99	0.97	0.92	0.99	0.99
0.98	0.99	0.99	0.85	0.81
0.6	0.99	0.99	0.91	0.89
0.94	0.99	0.97	0.74	0.94
0.88	0.98	0.98	0.84	0.96
0.96	1	1	0.97	0.87
0.92	0.88	0.89	0.95	0.96
0.9	0.98	0.99	0.97	0.96
0.8	1	0.95	0.96	0.89
0.95	0.99	0.99	0.94	0.94
0.96	0.14	1	0.97	0.75
0.75	0.97	0.9	0.88	0.93
0.87	0.96	0.93	0.95	0.9
0.88	0.9	1	0.27	0.93
	0.94		0.83	0.98
0.97	0.99	1	0.97	0.95
0.77	0.98	0.4	0.94	0.88
0.98	0.99	0.94	0.94	0.42
0.95	0.98	0.97	0.85	0.74
0.98	0.96	0.99	0.95	0.89
0.96	0.91	0.94	0.93	0.94
0.75	0.85	0.98	0.75	0.95
0.93	0.95	0.9	0.95	0.87
0.88	0.96	0.81	0.86	0.88
0.9	0.97	0.97	0.54	0.75
0.97	0.98	0.92	0.97	0.94
0.7	0.91	0.94	0.89	0.49
0.99	0.87		0.94	0.91
0.92	0.97	0.93	0.58	0.51
0.85	0.93	0.97	0.6	0.62
0.81	0.88	0.85	0.95	0.85

Millipore

0.84	0.97	0.95	0.83	0.82
0.91	0.98	0.9	0.64	0.58
0.78	0.93	0.87	0.65	0.82
0.81	0.96	0.51	0.82	0.9
0.3	0.95		0.38	0.65
0.94	0.99	0.99	0.57	0.57
0.99	0.89			
0.91	0.95	0.84	0.29	0.64
0.76	0.96	1		0.87
0.95	0.95	0.97	0.85	0.78
		0.89		
1	0.81			0.63
0.9	0.97	0.88	0.84	0.9
0.68	0.95	0.98	0.46	0.81
0.9	0.96	0.96	0.91	0.86
0.87	0.94	0.94	0.76	0.87
0.89	1	1	0.96	0.63
0.77	0.99	0.96	0.91	0.94
0.96	0.93	0.92	0.65	0.73
0.98	1	1	0.99	0.99
0.84	0.9	0.99	0.4	0.85
0.98	0.98	0.99	0.96	0.92
0.93	0.86	0.81	0.89	0.38
0.89	0.92	0.96	0.61	0.38
0.86	0.9	0.98	0.83	0.59
0.92	0.97	0.88	0.92	0.84
0.91	0.72	0.88	0.93	0.82
0.94	0.97	0.96	0.88	0.83
0.98	0.99	0.98	0.9	0.78
0.98	0.99	0.99	0.96	0.97
0.81	0.98	1	0.94	0.9
0.99	0.99	0.96	0.96	0.97
0.91	0.87	0.88	0.74	0.88
0.85	0.88	0.99	0.69	0.08
0.77	0.89	0.93	0.89	0.93

Millipore

0.99	0.93	0.95	0.81	0.91
0.97	0.99	0.98	0.7	0.87
0.98	0.93	0.97	0.5	0.47
0.95	0.85	0.97	0.61	0.49
0.99	0.98	1	0.76	0.67
0.91	0.86	0.77	0.75	0.47
0.91	0.88	0.95	0.93	0.95
0.91	0.9	0.8	0.69	0.61
0.99	0.99	1	0.54	0.64
0.94	0.73	0.98	0.99	0.54
0.99	0.98	1	0.74	0.67
0.99	0.96	0.99	0.54	0.55
0.59	0.82	0.73	0.65	0.33
0.93	0.92	0.9	0.52	0.57
0.55	0.89	0.65	0.68	0.67
0.72	0.99	0.98	0.96	0.68
0.8	0.95	0.93	0.77	0.7
0.88	0.91	0.94	0.72	0.67
0.87	0.93	0.82	0.6	0.72
0.96	0.98	0.95	0.8	0.86
0.94	0.7	0.86	0.58	0.72
0.88	0.96	0.99	0.62	0.68
0.78	0.86	0.96	0.93	0.85
0.6	0.81	0.49		0.53

Millipore

T3edta_across_ICC	T3heparin_across_ICC	T0HEPARIN_VS_T0SERUM_RHO	T3edta_VS_T3heparin_RHO
0.58	0.81	0.3286	0.2909
0.94	0.96	0.6374	0.8307
0.95	0.98	0.8234	0.8262
0.93	0.96	0.7219	0.7332
0.67	0.84	0.3469	0.3993
0.98	1	0.9547	0.9493
0.9	0.95	0.7345	0.7819
0.94	0.94	0.4862	0.456
0.76	0.94	0.3039	0.4444
0.97	0.93	0.8297	0.8587
0.99	0.99	0.9388	0.8138
0.81	0.99	0.4196	0.6443
0.98	0.99	0.5002	0.6362
0.8	0.74	0.6079	0.3989
0.95	0.93	0.8397	0.847
0.44	0.95	-0.1183	0.5
0.85	0.99	0.2162	0.7751
0.7	0.98	0.6222	0.6807
0.91	1	0.4987	0.4455
1	0.25	0.3095	0.619
0.99	0.96	0.8654	0.8941
0.9	0.89	0.5858	0.3573
0.83	0.93	0.651	0.7504
0.87	0.97	0.5652	0.7103
0.94	0.96	0.2337	0.5953
0.85	0.97	0.4675	0.4107
0.86	0.94	0.4152	0.4216
0.94	0.62	0.7246	0.4461
0.91	0.91	0.4137	0.5516
0.86	0.98	0.5024	0.5195
0.95	0.99	0.7471	0.8848
0.73	0.92	0.5106	0.5927
0.81	0.92	0.5578	0.4681
0.69	0.99	0.5045	0.4272
0.8	0.64	0.5834	0.6219
0.94	0.65	0.3354	0.3225

Millipore

0.93	0.85	0.5369	0.7735
0.66	0.98	0.6079	0.5273
0.74	0.94	0.5437	0.6881
0.84	1	0.4947	0.5824
0.78	0.36	0.2	0.3214
		1	
0.91	0.94	0.8295	0.8921
0.81	0.99	0.9524	0.119
0.97	0.95	0.7744	0.7063
0.99	0.76	0.5429	-0.4643
0.85	0.97	0.8367	0.5026
0.68	1	1	0.8
0.82	0.98	0.7048	0.6949
0.8	0.99	0.5162	0.556
0.9	0.92	0.416	0.701
0.92	0.99	0.1636	0.7143
0.95	0.99	0.8701	0.8134
0.97	0.84	0.6808	0.9085
0.61	0.84	0.5551	0.3093
0.99	1	0.9633	0.9659
0.8	0.76	0.6953	0.3831
0.7	0.94	0.4197	0.6384
0.82	0.79	0.5881	0.4775
0.77	0.93	0.6958	0.5395
0.81	0.67	0.2963	0.4036
0.95	0.98	0.6726	0.821
0.65	0.98	0.4424	0.57
0.91	0.99	0.5432	0.7197
0.79	0.91	0.6664	0.8038
0.99	1	0.9007	0.8662
0.93	0.99	0.8234	0.8024
0.98	0.99	0.8959	0.9441
0.77	0.93	0.4747	0.495
0.28	0.92	0.5371	0.8632
0.79	0.96	0.2412	0.4974

Millipore

0.83	0.97	0.6706	0.7515
0.79	0.97	0.487	0.5999
0.41	0.83	0.2455	0.3845
0.49	0.64	0.1421	0.2796
0.48	0.92	0.5389	0.6023
0.59	0.94	0.5158	0.15
0.85	1	0.8459	0.8462
0.57	0.86	0.5742	0.5576
0.64	0.93	0.6286	0.6497
0.95	0.38	0.4351	0.3326
0.73	0.99	0.5871	0.6343
0.58	0.82	0.6384	0.5376
0.42	0.21	0.1091	0.086
0.38	0.95	0.3937	0.4636
0.57	0.97	0.4212	0.6296
0.93	0.93	0.6437	0.6138
0.91	0.99	0.61	0.7
0.89	0.97	0.4733	0.6491
0.89	0.99	0.5604	0.454
0.87	0.97	0.645	0.7636
0.81	0.84	0.5291	0.6766
0.95	1	0.8679	0.9364
0.64	0.98	0.0689	0.5
	0.51	0.4182	-0.2