

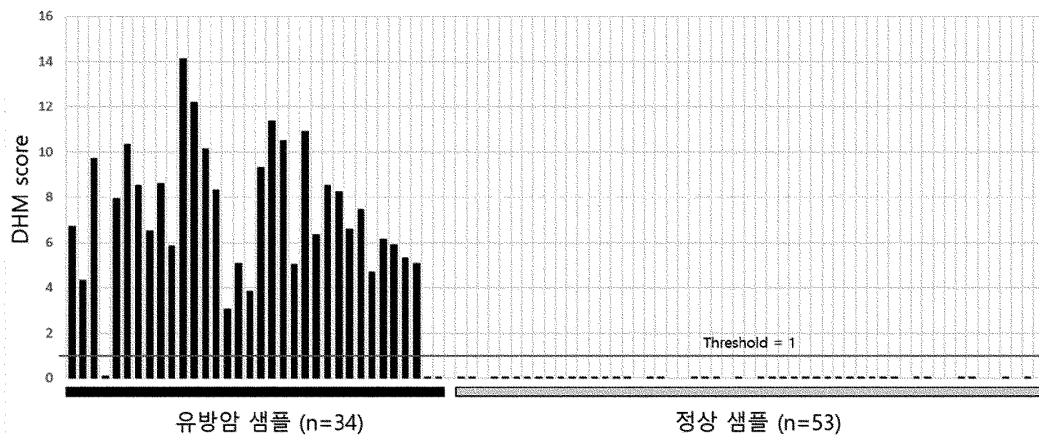
[51a]

유방암 샘플 (n=34)

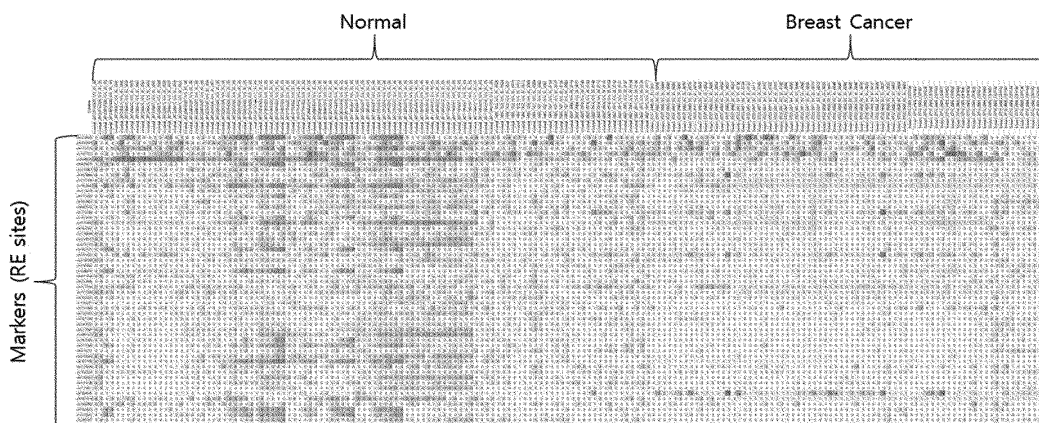
정상 샘플 (n=53)

arXiv	Normalized 0001	Normalized 0002	Normalized 0003	Normalized 0004	Normalized 0005	Normalized 0006	Normalized 0007	Normalized 0008	Normalized 0009	Normalized 0010	Normalized 0011	Normalized 0012	Normalized 0013	Normalized 0014	Normalized 0015	Normalized 0016	Normalized 0017	Normalized 0018	Normalized 0019	Normalized 0020	Normalized 0021	Normalized 0022	Normalized 0023	Normalized 0024	Normalized 0025	Normalized 0026	Normalized 0027	Normalized 0028	Normalized 0029	Normalized 0030
CG0000	0.421	22.20	34.15	11.12	27.24	20.18	4.9	8.22	23.81	8.14	9.15	17.10	1.18	9.18	9.12	10.1														
CG0001	0.913	19.18	11.13	18.12	19.22	16.14	8.9	8.16	16.33	18.9	6.21	13.13	10.1	10.12	11.21	5.6														
CG0002	0.913	19.18	11.13	18.12	19.22	16.14	8.9	8.16	16.33	18.9	6.21	13.13	10.1	10.12	11.21	5.6														
CG0003	0.913	19.18	11.13	18.12	19.22	16.14	8.9	8.16	16.33	18.9	6.21	13.13	10.1	10.12	11.21	5.6														
CG0004	0.913	12.12	12.15	14.10	27.16	14.12	5.8	5.13	17.14	4.16	5.14	18.1	8.1	7.13	8.30	11.1														
CG0005	0.913	19.17	15.1	8.10	13.28	13.17	7.9	3.19	18.17	3.16	7.12	13.15	10.1	9.12	13.1															
CG0006	1.3	6.19	7.20	7.1	5.12	27.22	10.12	5.7	7.14	12.15	8.18	8.11	6.91	1.92	1.92															
CG0007	0.913	19.18	12.1	7.13	12.31	19.16	4.4	6.115	20.16	4.12	4.152	9.12	13.1	7.9	6.151															
CG0008	0.9	8.22	13.12	15.12	11.24	20.14	14.7	7.11	5.16	18.11	7.14	6.14	14.11	6.1	7.111	7.6														
CG0009	1.3	7.9	9.25	13.7	1.15	21.14	5.9	7.14	17.7	6.18	10.13	11.1	8.18	8.9	7.8	9.6														
CG0010	0.913	19.18	6.11	10.1	20.14	14.7	5.9	5.11	19.15	8.1	7.15	14.17	1.1	8.15	14.17	1.1														
CG0011	1.2	4.20	15.14	7.14	13.10	26.18	14.8	9.9	5.4	13.17	6.13	7.12	10.12	10.6	9.9	12.7														
CG0012	1.1	7.10	5.11	16.9	9.12	5.10	16.19	14.4	4.6	10.16	17.9	2.05	4.14	10.12	10.6	10.9	10.0													
CG0013	0.913	19.18	12.1	5.11	8.15	19.14	10.1	10.1	6.11	11.81	6.10	7.15	12.13	10.1	10.1	10.1														
CG0014	0.913	19.18	13.1	10.1	17.1	6.18	15.1	4.4	8.112	16.14	7.14	7.10	10.10	6.10	7.7	10.0														
CG0015	0.913	1.81	1.11	1.11	7.14	4.9	6.151	9.3	4.7	14.16	16.15	7.13	13.10	7.6	11.7	6.7														
CG0016	0.913	19.18	12.1	1.14	7.1	7.12	12.1	7.1	10.15	19.1	7.15	10.15	7.15	10.15	10.15	10.15														
CG0017	0.913	5.8	13.14	5.1	6.10	15.10	10.1	4.4	4.14	14.12	6.12	7.13	11.9	9.12	6.6	5.8														
CG0018	0.913	8.16	6.14	6.14	6.14	10.10	10.1	5.4	4.19	10.14	5.9	5.9	11.15	1.5	7.7	7.5														
CG0019	0.913	8.16	6.14	6.14	6.14	10.10	10.1	5.4	4.19	10.14	5.9	5.9	11.15	1.5	7.7	7.5														
CG0020	0.913	4.31	9.11	5.8	5.8	4.16	13.9	8.9	4.6	10.15	7.6	6.32	6.10	12.5	5.11	7.1	4.9													
CG0021	0.913	7.7	7.50	1.9	8.18	11.1	7.2	2.6	8.9	10.14	6.11	4.9	1.8	7.6	6.7	1.1														
CG0022	1.5	1.53	0.81	6.1	6.1	5.15	19.10	10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.1														
CG0023	0.913	5.8	13.14	7.7	4.7	14.15	10.1	2.6	4.6	10.10	10.1	8.7	7.8	1.15	5.1	7.7	6.1													
CG0024	0.913	7.4	13.1	7.9	9.9	8.6	4.37	12.11	8.5	3.2	7.35	12.4	4.4	7.10	5.4	4.6	5.9													
CG0025	0.913	4.7	6.1	7.9	7.9	1.9	5.1	9.10	1.5	5.1	12.10	10.1	4.4	7.10	5.4	4.6	5.9													
CG0026	0.913	6.9	9.9	9.9	5.1	12.10	10.1	8.6	4.6	10.11	10.1	10.1	10.1	10.1	10.1	10.1														
CG0027	0.913	6.3	13.2	8.10	3.9	7.12	15.10	10.7	2.5	3.10	12.1	5.9	4.10	1.3	3.5	4.4														
CG0028	0.913	5.10	6.16	9.8	10.1	8.15	4.4	4.4	4.10	12.1	5.13	3.9	7.8	7.8	1.8	4.6	5.9													
CG0029	0.913	4.9	8.7	7.10	4.35	8.7	1.1	4.8	10.10	10.1	4.10	5.1	4.10	5.1	4.10	5.1														
CG0030	0.913	6.4	5.1	10.1	5.1	8.12	14.9	3.2	5.9	10.14	8.4	13.1	7.8	5.6	4.8	6.5														
CG0031	0.913	5.13	6.9	8.9	6.7	7.16	13.11	9.3	4.6	8.10	10.1	6.12	6.9	5.1	4.6	5.3														
CG0032	0.913	4.9	10.1	7.7	7.14	12.1	5.9	2.4	9.9	8.1	4.6	5.10	10.1	6.1	4.6	5.3														
CG0033	0.913	4.3	1.12	1.32	10.1	6.7	2.3	4.9	9.13	7.11	4.6	8.11	5.1	5.5	5.8	5.8														
CG0034	0.913	5.7	6.11	6.9	8.6	6.10	11.8	6.1	5.5	13.10	10.1	7.8	11.7	7.4	4.7	6.5														
CG0035	0.913	7.14	6.9	8.1	6.12	10.1	8.3	3.1	9.110	10.1	6.10	5.9	6.4	3.1	4.5	5.5														
CG0036	1.2	1.3	5.1	5.7	6.3	10.11	14.6	1.4	4.6	6.12	9.5	6.5	7.10	7.1	4.9	7.5														
CG0037	0.913	4.1	7.1	3.1	8.1	10.1	12.1	14.6	2.5	6.18	10.1	5.14	4.9	7.1	4.1	4.6	3.5													
CG0038	0.913	4.4	7.12	4.11	5.11	10.1	10.1	6.2	3.9	10.12	7.12	7.14	9.8	8.10	4.8	4.8														
CG0039	0.913	7.7	8.10	15.11	6.1	5.11	1.8	2.4	4.9	10.13	10.1	10.1	10.1	10.1	10.1	10.1														
CG0040	0.913	8.10	8.12	6.1	5.10	10.10	10.1	3.6	5.10	8.11	5.10	5.6	7.8	5.4	4.8	5.5														
CG0041	0.913	4.13	7.18	4.9	6.13	15.1	5.2	4.8	10.11	4.13	4.8	6.8	6.1	4.5	4.4	5.4														
CG0042	0.913	6.16	6.16	8.15	12.1	10.10	10.1	3.1	7.12	15.11	7.11	9.9	7.1	6.1	5.1	5.1														
CG0043	0.913	6.10	10.1	5.1	6.14	3.9	4.4	7.1	7.10	12.1	7.10	5.7	5.9	1.5	5.5	5.4														
CG0044	0.913	4.21	1.8	6.9	6.1	2.23	8.12	7.2	4.4	7.10	12.1	5.16	4.8	9.1	5.4	5.9														
CG0045	0.913	4.8	7.7	1.2	2.16	10.11	10.1	2.6	6.9	8.12	4.4	6.1	6.1	6.1	6.1	6.1														
CG0046	0.913	7.4	1.10	6.14	10.1	9.17	11.1	9.0	3.6	6.10	7.4	4.8	7.6	7.6	4.4	5.2														
CG0047	0.913	6.9	8.1	6.7	4.8	10.11	10.1	3.8	6.1	13.10	10.1	4.8	7.7	3.9	8.3	9.3														
CG0048	0.913	8.7	7.10	6.7	7.13	10.1	10.1	3.9	10.13	10.1	4.8	7.7	3.9	8.3	9.3	9.3														
CG0049	0.913	4.1	4.12	5.9	3.11	12.9	4.4	3.5	3.5	11.11	4.15	5.5	7.5	7.5	7.5	7.5														
CG0050	0.913	6.4	8.1	6.7	7.8	11.13	10.1	4.4	9.8	10.1	5.12	3.7	7.5	1.4	4.4	3.9														
CG0051	0.913	4.1	5.9	7.8	6.11	12.1	4.6	2.5	4.9	10.1	10.1	4.8	7.7	3.9	8.3	9.3														
CG0052	0.913	4.9	7.10	9.5	6.13	12.12	4.6	2.4	4.7	10.1	7.3	6.6	4.4	4.4	4.4	4.4														
CG0053	0.913	7.3	1.10	7.8	6.9	8.16	11.1	7.8	4.6	6.11	6.4	4.9	7.10	4.4	3.2	7.3	3.5													
CG0054	0.913	2.9	7.7	4.7	7.7	8.10	10.1	6.7	4.11	9.18	5.32	3.11	1.1	1.1	1.1	1.1														
CG0055	0.913	4.1	6.1	6.1	5.1	12.12	8.1	2.1	7.1	10.1	8.1	4.8	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	7.1	

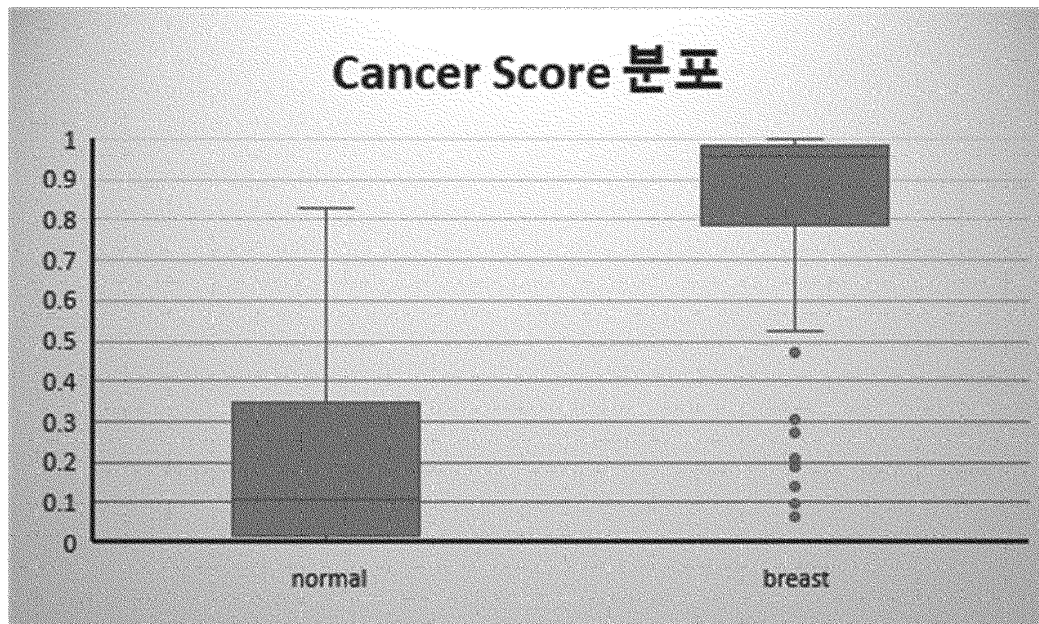
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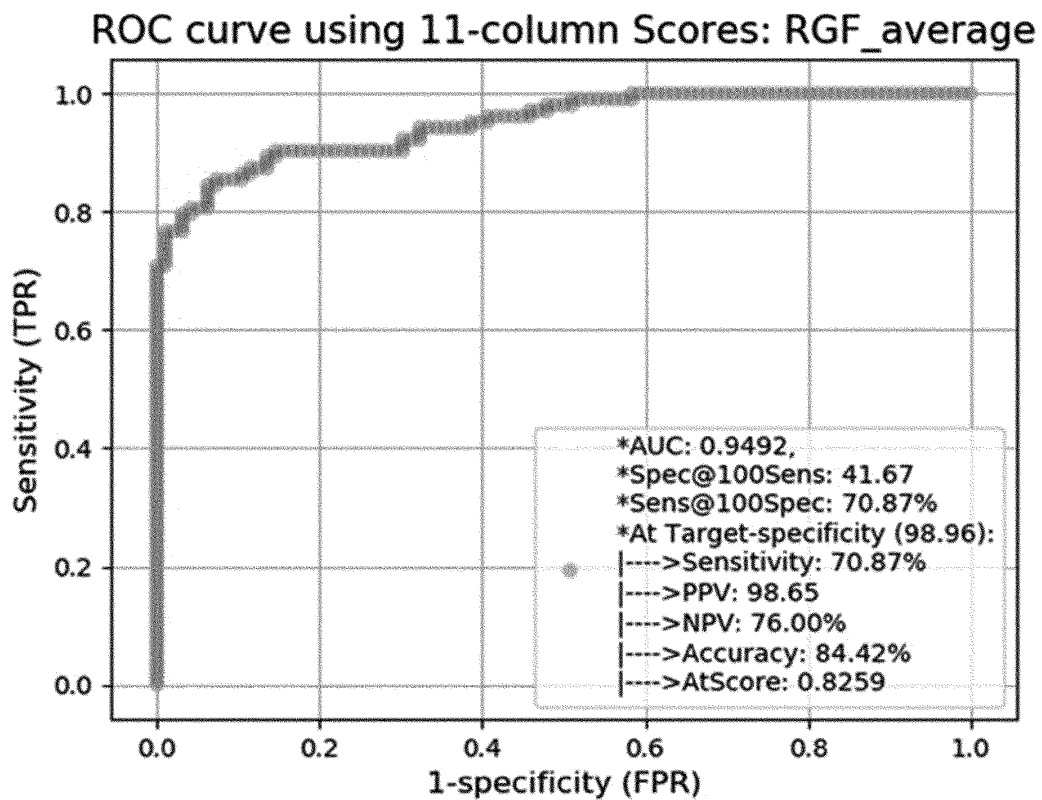
[E2a]



[도2b]



[도2c]



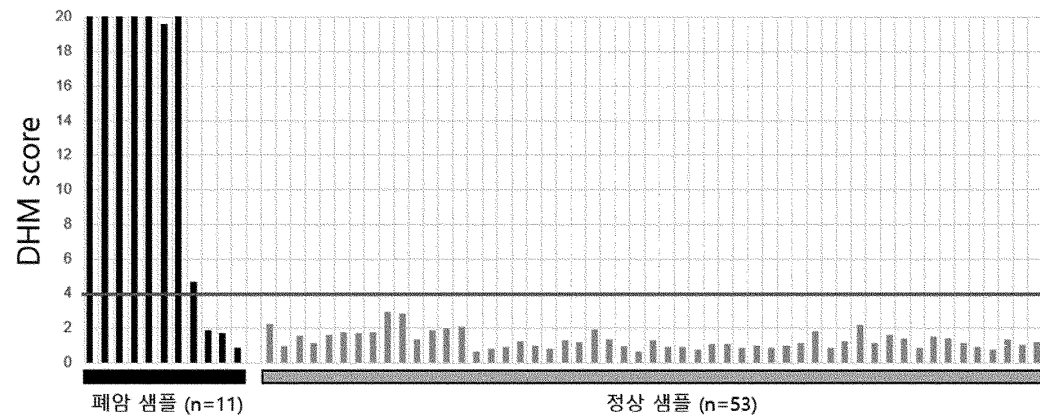
[도3a]

폐암 샘플 (n=11)

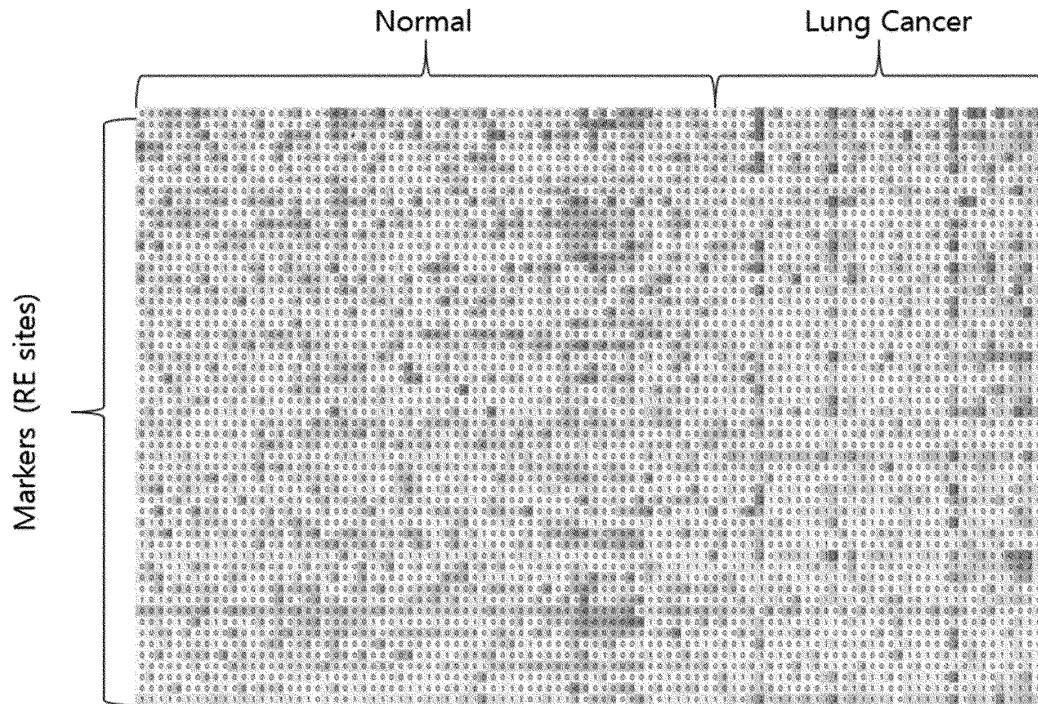
정상 샘플 (n=53)

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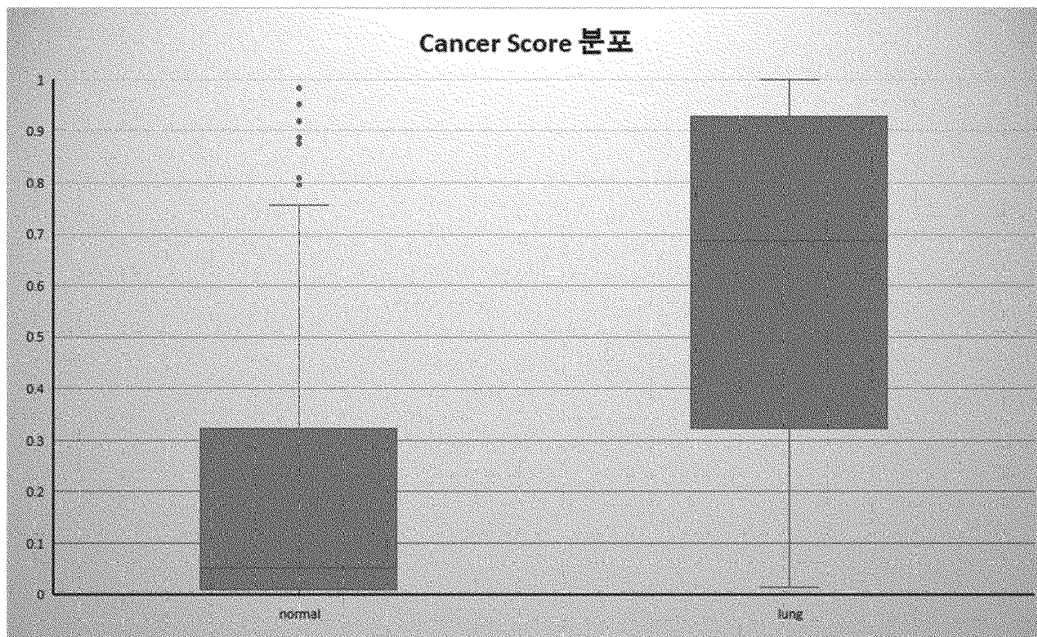
[도3b]



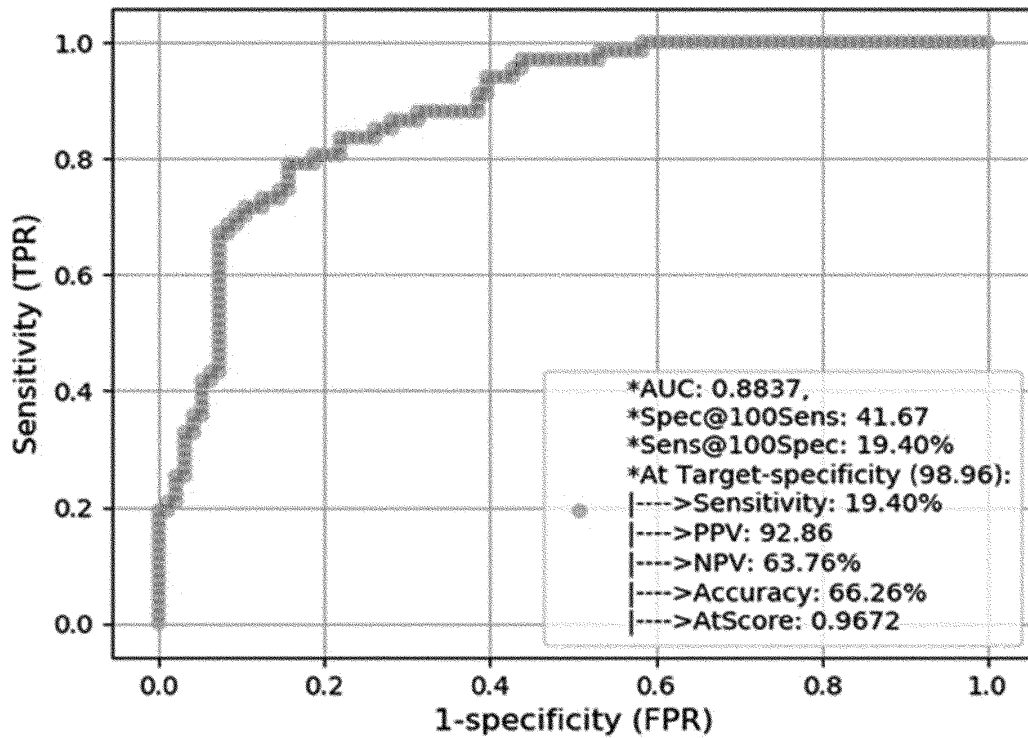
[도4a]



[도4b]



[도4c]



[도4d]

