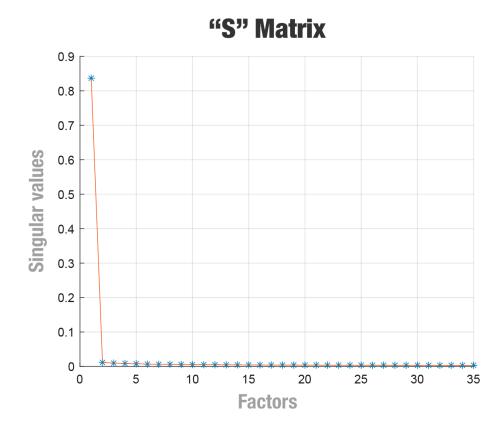
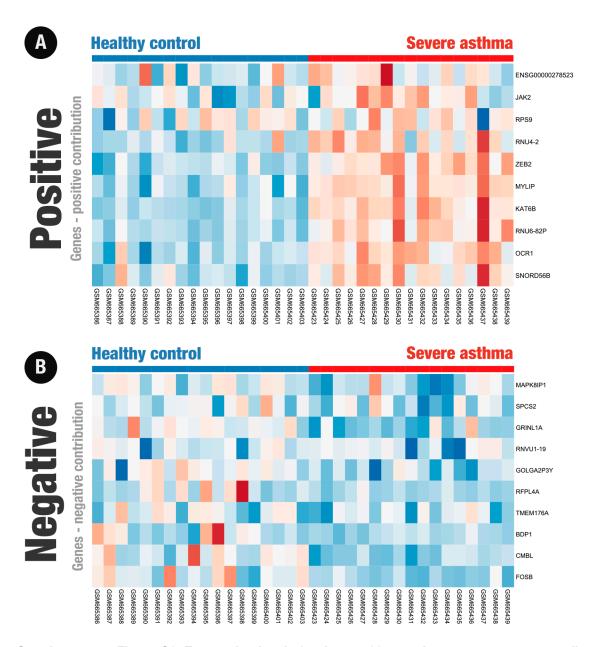
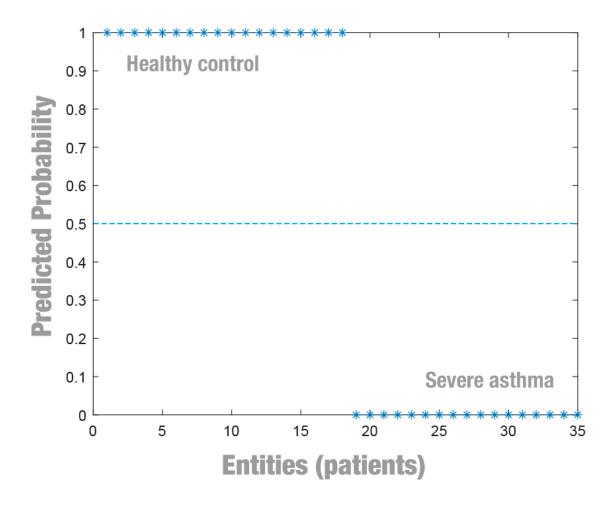
Supplementary material for "Mapping Relevant Genes in Severe Asthma: A Computational Strategy for Biomarker Discovery"



Supplementary Figure S1. Importance of singular values by total factors (principal components).



Supplementary Figure S2. Expression levels for the top 20 most important genes according to the weight distribution (α) derived from the logistic regression model. Values were normalized by row. Details are available at Supplementary Table S2 and S3.



Supplementary Figure S3. Classification probabilities obtained by Logistic Regression. Scores close to 1: healthy control; scores close to 0: severe asthma.

Supplementary Table S1. Expression levels of the top 10 most important genes by the Information Gain algorithm (HC: Healthy control; SA: Severe asthma)

	Clas											
#	S	ID	7944525	8005829	8157727	7907531	7968295	8117018	8144512	7928489	8103620	8151411
1	HC	GSM665386	6.89	4.85	6.06	6.08	3.88	4.68	2.75	6.79	2.44	7.77
2	HC	GSM665387	6.74	4.25	5.86	6.34	3.32	4.03	2.84	6.86	2.57	7.26
3	HC	GSM665388	6.94	5.51	6.04	6.58	4.40	5.42	2.65	7.45	2.44	7.68
4	HC	GSM665389	6.85	4.79	5.47	6.26	3.71	4.49	2.63	6.92	2.44	7.67
5	HC	GSM665390	6.77	4.35	5.65	6.30	3.64	3.38	2.74	7.28	2.53	7.28
6	HC	GSM665391	6.82	5.20	6.19	6.83	3.84	5.50	2.62	7.16	2.68	7.63
7	HC	GSM665392	6.27	4.90	5.69	5.95	3.80	4.83	2.64	7.09	2.47	7.02
8	HC	GSM665393	6.39	5.28	6.35	6.70	3.73	4.72	2.62	7.04	2.51	7.62
9	HC	GSM665394	6.57	4.04	5.97	6.05	3.32	3.79	2.70	6.87	2.32	7.47
10	HC	GSM665395	6.41	5.04	6.03	5.64	3.60	4.59	2.64	6.77	2.34	7.33
11	HC	GSM665396	6.53	5.02	5.16	6.02	3.45	4.73	2.64	6.84	2.60	7.59
12	HC	GSM665397	6.95	5.10	6.23	6.72	4.09	5.92	2.73	7.52	2.76	7.54
13	HC	GSM665398	6.38	4.00	5.75	6.11	3.58	4.39	2.72	7.05	2.49	7.38
14	HC	GSM665399	6.62	5.13	6.04	6.72	3.91	4.93	2.65	7.27	2.51	7.64
15	HC	GSM665400	6.78	4.86	6.10	6.29	4.18	5.59	2.66	7.69	2.49	7.53
16	HC	GSM665401	6.12	4.56	5.90	5.98	3.38	3.62	2.74	7.04	2.53	7.65
17	HC	GSM665402	6.63	4.96	6.15	6.50	3.92	5.54	2.78	7.10	2.49	7.51
18	HC	GSM665403	6.57	4.35	5.71	6.06	3.25	3.82	2.57	6.64	2.74	7.68
19	SA	GSM665423	7.36	5.82	6.90	7.32	4.28	5.92	2.93	8.26	2.40	7.93
20	SA	GSM665424	6.47	5.42	6.36	6.73	4.49	6.09	2.84	8.06	2.43	7.72
21	SA	GSM665425	7.03	5.55	6.84	7.11	5.19	6.57	2.98	8.20	2.33	7.76
22	SA	GSM665426	7.02	5.39	6.26	6.97	4.72	6.52	2.85	7.84	2.38	7.69
23	SA	GSM665427	7.00	6.04	6.94	7.28	5.13	6.34	2.87	8.51	2.42	8.09
24	SA	GSM665428	7.11	6.04	6.76	6.76	4.60	6.24	2.82	8.26	2.35	7.86
25	SA	GSM665429	7.13	5.87	6.91	6.84	4.40	5.84	2.72	8.21	2.30	8.04
26	SA	GSM665430	7.68	6.73	7.49	7.83	6.00	7.64	2.96	8.94	2.40	8.19
27	SA	GSM665431	7.25	4.72	6.07	6.60	4.17	5.51	2.93	7.52	2.28	7.93
28	SA	GSM665432	7.35	6.68	7.19	7.71	5.35	7.16	2.96	8.90	2.42	7.88
29	SA	GSM665433	7.32	5.36	6.29	7.00	5.03	6.00	3.08	8.57	2.41	7.56
30	SA	GSM665434	7.04	5.83	6.91	7.32	5.04	6.00	2.90	8.23	2.23	7.65
31	SA	GSM665435	7.12	5.90	7.02	7.22	4.44	6.06	2.93	7.86	2.53	8.00
32	SA	GSM665436	7.06	5.56	7.09	7.17	4.46	5.87	2.79	7.71	2.38	8.07
33	SA	GSM665437	7.73	7.05	7.70	7.88	6.29	7.70	3.12	9.51	2.43	8.62
34	SA	GSM665438	7.21	5.57	6.67	7.08	4.59	6.46	3.02	7.97	2.42	7.86
35	SA	GSM665439	7.44	6.73	7.13	7.64	5.58	6.40	2.92	8.02	2.38	7.95

Genes: 7944525: RNU6-1123P; 8005829: PYY2; 8157727: GPR21; 7907531: GPR52; 7968295: RNU6-82P; 8117018: RNU6-522P; 8144512: RNU6-682P; 7928489: KAT6B; 8103620: RNU6-1336P; 8151411: TCEB1.

Supplementary table S2. Expression levels of the top 10 genes that contribute most negatively (Healthy controls class) to the model, according to the logistic regression model

class	Negative												
	gene	8137264	7983890	8110971	7904967	8106025	7985431	8031570	8007848	8029693	7914180		
Healthy control	GSM66538 6	6.23	7.20	6.39	9.55	4.30	5.84	3.02	5.23	8.45	4.96		
Healthy control	GSM66538	7.64	6.63	7.03	9.43	3.72	6.95	3.63	6.06	9.03	3.87		
Healthy control	GSM66538 8	9.29	6.54	5.61	9.02	3.48	4.89	3.34	6.18	7.86	4.45		
Healthy control	GSM66538	7.24	8.23	5.66	9.32	2.16	6.82	3.44	5.98	7.94	4.89		
Healthy control	GSM66539 0	7.60	6.61	6.10	6.58	2.55	7.01	4.49	4.76	7.83	4.19		
Healthy control	GSM66539	8.77	7.10	5.95	8.24	2.17	7.30	4.99	5.80	6.92	3.57		
Healthy control	GSM66539 2	8.32	7.45	5.23	9.47	2.55	6.31	3.43	6.09	9.84	4.71		
Healthy control	GSM66539 3	6.21	7.18	6.04	8.83	2.32	7.49	4.86	4.31	7.33	4.80		
Healthy control	GSM66539 4	8.42	7.06	7.73	8.45	1.76	6.97	3.31	5.49	7.98	5.01		
Healthy control	GSM66539 5	8.48	7.37	5.94	8.64	4.26	6.77	5.60	5.17	7.45	5.03		
Healthy control	GSM66539 6	8.18	7.23	6.77	8.75	5.44	6.29	3.66	6.25	8.88	3.00		
Healthy control	GSM66539	7.37	6.74	6.31	9.30	2.73	6.61	4.85	6.29	9.73	5.03		
Healthy control	GSM66539 8	6.38	6.81	5.69	7.08	3.42	7.07	6.93	4.93	6.80	5.61		
Healthy control	GSM66539 9	5.91	7.22	5.64	8.66	3.24	6.64	3.76	5.28	6.51	3.95		
Healthy control	GSM66540 0	7.78	7.04	6.53	9.49	2.51	6.99	3.06	6.17	8.22	6.21		
Healthy control	GSM66540	8.44	6.93	6.30	9.69	2.35	6.58	3.46	6.10	7.89	4.76		
Healthy control	GSM66540 2	8.46	7.56	6.32	8.38	2.75	6.75	3.31	5.08	7.86	3.35		
Healthy control	GSM66540 3	6.15	6.37	6.59	8.96	2.22	6.00	3.97	5.18	8.77	4.85		
Severe asthma	GSM66542 3	6.20	5.84	5.01	9.03	2.44	6.46	3.56	4.35	7.34	3.94		
Severe asthma	GSM66542 4	5.74	7.15	5.30	9.30	2.37	5.75	2.92	3.98	7.28	3.23		
Severe asthma	GSM66542 5	8.01	5.92	5.45	9.02	1.99	6.53	3.33	5.32	8.05	4.92		
Severe asthma	GSM66542 6	7.60	6.17	5.47	9.20	2.38	5.80	3.63	5.09	7.42	5.13		
Severe	GSM66542 7	8.00	6.28	5.95	9.27	2.10	6.39	4.38	5.11	7.11	3.45		
Severe asthma	GSM66542 8	6.64	6.51	5.36	8.90	2.55	4.85	3.23	7.34	6.93	5.85		
Severe	GSM66542	8.09	6.15	5.33	9.42	2.68	6.08	3.27	5.20	7.12	5.00		
Severe	9 GSM66543	7.86	6.43	5.63	8.81	2.88	6.96	3.63	5.46	7.68	3.46		
Severe	0 GSM66543	5.74	5.99	5.02	6.42	2.62	6.74	3.17	4.70	6.74	4.25		
Severe	GSM66543	8.25	5.91	5.46	8.30	2.91	6.24	3.35	4.08	6.69	2.57		
asthma	2												

Severe	GSM66543	6.57	7.40	5.37	9.21	2.40	5.48	3.43	3.60	7.31	3.35
asthma	3										
Severe	GSM66543	7.77	6.14	6.02	6.84	2.43	6.53	3.99	3.87	6.91	2.96
asthma	4										
Severe	GSM66543	7.88	6.61	5.96	6.55	2.56	5.58	3.35	4.62	6.88	5.27
asthma	5										
Severe	GSM66543	8.69	7.85	5.94	9.36	3.15	5.32	3.26	5.77	7.07	3.53
asthma	6										
Severe	GSM66543	7.88	6.47	5.64	9.53	2.50	6.32	3.51	6.02	7.03	4.93
asthma	7										
Severe	GSM66543	6.02	6.35	5.77	9.56	2.46	6.31	3.17	5.80	8.26	4.64
asthma	8										
Severe	GSM66543	6.76	6.40	5.81	8.52	2.30	6.55	3.30	4.93	6.62	3.43
asthma	9										

Supplementary table S3. The top 10 genes that contribute most positively (Severe asthma class) to the model, according to the logistic regression model

		Positive										
		811701	790886	796702	792848	805258	815421	796829	803115	805563 797545		
class	gene	8	1	8	9	1	1	5	2	9	3	
Healthy	GSM66											
control	5386	4.68	6.95	6.01	6.79	4.34	7.01	3.88	3.96	5.72	3.73	
Healthy	GSM66											
control	5387	4.03	6.16	6.20	6.86	4.14	6.44	3.32	2.77	5.82	4.00	
Healthy	GSM66											
control	5388	5.42	8.51	6.21	7.45	3.54	7.04	4.40	5.10	6.92	6.02	
Healthy	GSM66											
control	5389	4.49	6.95	6.86	6.92	3.71	5.96	3.71	3.99	6.30	4.78	
Healthy	GSM66	0.00	F 77	0.00	7.00	F 00	0.44	0.04	F 70	F 70	4.00	
control	5390	3.38	5.77	6.23	7.28	5.96	6.14	3.64	5.72	5.76	4.06	
Healthy	GSM66	E E0	7.04	7.04	7 16	2.07	6.61	2 04	4 22	6.45	4.00	
control	5391	5.50	7.04	7.04	7.16	3.07	6.61	3.84	4.33	6.45	4.82	
Healthy	GSM66	4.83	6.87	6 44	7.09	4.99	7.07	2 00	2.42	6.67	E 25	
control Healthy	5392 GSM66	4.03	0.07	6.44	7.09	4.99	7.07	3.80	3.42	6.67	5.35	
control	5393	4.72	7.42	5.45	7.04	2.80	6.49	3.73	5.58	6.50	4.21	
Healthy	GSM66	4.72	7.42	3.43	7.04	2.00	0.49	3.73	5.56	0.50	4.21	
control	5394	3.79	6.39	6.08	6.87	4.69	5.53	3.32	4.41	6.40	3.84	
Healthy	GSM66	3.73	0.55	0.00	0.07	4.03	3.33	3.32	7.71	0.40	3.04	
control	5395	4.59	7.51	5.78	6.77	3.45	6.97	3.60	5.84	6.39	4.58	
Healthy	GSM66	7.00	7.01	3.70	0.11	0.40	0.57	3.00	3.04	0.00	7.50	
control	5396	4.73	6.75	6.08	6.84	3.89	4.98	3.45	5.26	6.80	4.69	
Healthy	GSM66	4.73	0.73	0.00	0.04	3.03	7.50	0.40	3.20	0.00	7.00	
control	5397	5.92	7.35	7.62	7.52	3.96	5.11	4.09	5.35	6.59	4.44	
Healthy	GSM66	0.02	7.00	7.02	7.02	0.00	0.11	4.00	0.00	0.00	7.77	
control	5398	4.39	6.84	6.41	7.05	4.12	5.98	3.58	4.13	6.07	3.58	
Healthy	GSM66	1.00	0.01	0.11	7.00	1.12	0.00	0.00	1.10	0.01	0.00	
control	5399	4.93	7.63	6.39	7.27	3.17	5.79	3.91	5.67	7.00	4.91	
Healthy	GSM66											
control	5400	5.59	6.90	6.26	7.69	4.39	7.16	4.18	5.12	6.14	4.14	
Healthy	GSM66											
control	5401	3.62	7.21	8.48	7.04	5.52	5.55	3.38	5.10	6.10	4.34	
Healthy	GSM66											
control	5402	5.54	7.36	6.14	7.10	3.93	6.86	3.92	5.44	6.45	4.22	
Healthy	GSM66											
control	5403	3.82	7.17	6.19	6.64	3.72	6.64	3.25	3.78	6.08	4.31	
Severe	GSM66											
asthma	5423	5.92	8.19	8.26	8.26	5.43	5.23	4.28	5.00	7.03	5.32	
Severe	GSM66											
asthma	5424	6.09	8.07	7.98	8.06	5.23	7.08	4.49	4.45	6.68	4.77	
Severe	GSM66											
asthma	5425	6.57	8.36	8.80	8.20	4.33	6.66	5.19	5.65	7.00	5.95	
Severe	GSM66											
asthma	5426	6.52	7.71	7.35	7.84	4.49	6.67	4.72	4.10	7.29	5.25	
Severe	GSM66											
asthma	5427	6.34	9.20	8.38	8.51	4.78	8.10	5.13	5.00	7.17	6.31	
Severe	GSM66											
asthma	5428	6.24	8.41	8.29	8.26	4.84	7.43	4.60	6.44	7.12	6.14	
Severe	GSM66											
asthma	5429	5.84	8.11	7.23	8.21	6.60	7.90	4.40	4.81	7.86	5.29	
Severe	GSM66						2.25					
asthma	5430	7.64	9.05	8.28	8.94	3.89	6.89	6.00	5.91	7.95	6.39	
Severe	GSM66											
asthma	5431	5.51	8.88	5.92	7.52	4.47	7.40	4.17	5.84	6.75	4.75	
Severe	GSM66	- 46			0.00					- 00		
asthma	5432	7.16	8.81	7.98	8.90	3.84	7.77	5.35	4.43	7.86	5.65	

Severe	GSM66										
asthma	5433	6.00	7.55	6.71	8.57	4.16	6.67	5.03	5.39	7.09	5.56
Severe	GSM66										
asthma	5434	6.00	7.67	7.85	8.23	5.04	6.89	5.04	5.79	7.39	4.74
Severe	GSM66										
asthma	5435	6.06	8.34	6.64	7.86	3.85	6.80	4.44	4.84	7.78	5.54
Severe	GSM66										
asthma	5436	5.87	8.56	7.66	7.71	4.90	7.75	4.46	5.48	7.31	4.79
Severe	GSM66										
asthma	5437	7.70	9.19	9.48	9.51	4.71	6.15	6.29	2.55	8.13	7.08
Severe	GSM66										
asthma	5438	6.46	8.85	7.88	7.97	3.97	5.73	4.59	5.14	6.90	5.69
Severe	GSM66										
asthma	5439	6.40	7.50	6.69	8.02	3.51	5.99	5.58	5.33	7.78	5.17