

Table S3. The entire list of consensus prognostic genes identified by the revolving sliding window approach.

NUMBER	GENE NAME	ACCURACY COUNT
1	LSP1	2
2	ACSBG2	3
3	POSTN	3
4	MMP1	3
5	HOXC8	3
6	COL3A1	4
7	PDZRN3	5
8	WNT2	5
9	CEND1	5
10	FCRL2	6
11	SLC19A1	6
12	PDGFRL	6
13	KCTD3	6
14	SNUPN	7
15	PNOC	7
16	ISG20	8
17	C1S	9
18	NEUROG1	10
19	YBX2	11
20	CCL2	11
21	CSTA	12
22	ITGB7	12
23	PRDM1	12
24	FAM46C	12
25	CCDC88C	13
26	FMO3	14
27	CD83	14
28	METTL2B	14
29	IGL@	14
30	KIAA0125	16
31	C2CD2L	16
32	IGJ	17
33	RHOH	18
34	GUSBP11	19
35	DCN	19
36	INPPL1	22
37	IGHM	23
38	MXI1	23
39	PIM2	24
40	ITM2A	25
41	IGKV4-1	25
42	IGLV3-25	26
43	LAX1	26
44	DTNB	28

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45	MZB1	29
46	IGLL3P	30
47	IGKC	31
48	POU2AF1	33
49	IGHG1	34
50	TNFRSF17	34
51	IGHD	34
52	VPREB3	34
53	CPA3	36
54	GM2A	38
55	IGLJ3	40
56	IGLV1-40	41
57	IGLV3-19	42
58	CD27	47
59	CD79A	48