

Name	Chromosome	Genomic coordinates	SVM score	Cloning	Conservation			Closest miRNA	Other predictions
HP-1	chr1	40887341-40887436,+	3.02	-	-	-	-	hsa-mir-30e	
HP-2	chr1	40898684-40898762,+	0.44	-	-	-	-	hsa-mir-30c-1	
HP-3	chr1	65232147-65232234,-	2.72	-	-	-	-	hsa-mir-101-1	
HP-4	chr1	98222840-98222917,-	0.48	-	m r	m r	r	hsa-mir-137	B648
HP-5	chr1	153211595-153211689,-	0.95	-	-	-	-	hsa-mir-9-1	
HP-6	chr2	136262939-136263027,+	0.06	-	-	-	-	hsa-mir-128a	
HP-7	chr2	176831714-176831785,+	0.04	-	-	-	-	hsa-mir-10b	
HP-8	chr3	37982219-37982304,+	0.46	-	-	-	-	hsa-mir-26a-1	
HP-9	chr3	161610634-161610741,+	0.10	-	-	-	-	hsa-mir-16-2	
HP-10	chr5	54503853-54503934,-	2.09	-	-	-	-	hsa-mir-449	
HP-11	chr5	148785556-148785635,+	1.21	-	-	-	-	hsa-mir-143	
HP-12	chr5	179377228-179377312,-	0.11	-	-	-	-	hsa-mir-340	
HP-13	chr6	52127783-52127846,+	0.44	-	-	-	-	hsa-mir-133b	
HP-14	chr6	72141346-72141459,-	0.74	-	m r	r	-	hsa-mir-30c-2	
HP-15	chr7	129017934-129018046,-	4.42	-	m	m	-	hsa-mir-183	
HP-16	chr7	129722221-129722300,+	0.58	-	m	-	-	hsa-mir-335	
HP-17	chr8	141804157-141804256,-	2.00	-	-	-	-	hsa-mir-151	
HP-18	chr8	141813107-141813217,-	0.09	-	-	-	-	hsa-mir-151	
HP-19	chr9	120085235-120085329,-	0.50	-	-	-	-	hsa-mir-147	
HP-20	chr11	553994-554064,-	0.31	-	-	-	-	hsa-mir-210	
HP-21	chr11	559747-559827,-	0.82	-	-	-	-	hsa-mir-210	
HP-22	chr11	74715663-74715760,-	1.78	-	m	m	-	hsa-mir-326	
HP-23	chr11	74716123-74716211,-	2.05	-	-	-	-	hsa-mir-326	
HP-24	chr11	74716486-74716607,-	0.44	-	m r	-	-	hsa-mir-326	
HP-25	chr11	121485215-121485296,-	2.41	-	-	-	-	hsa-mir-125b-1	
HP-26	chr11	121528013-121528085,-	0.47	-	-	-	-	hsa-mir-100	
HP-27	chr12	52668365-52668434,+	0.85	-	-	-	-	hsa-mir-196a-2	
HP-28	chr13	49519429-49519510,-	0.81	-	-	-	-	hsa-mir-16-1	
HP-29	chr14	99636013-99636131,+	0.85	-	-	-	-	hsa-mir-342	
HP-30	chr14	100405160-100405230,+	0.12	h	m r	m r	-	hsa-mir-337	B202, X52
HP-31	chr14	100420573-100420666,+	1.27	h	m	m	-	hsa-mir-136	
HP-32	chr14	100447566-100447644,+	0.42	-	-	-	-	hsa-mir-370	
HP-33	chr14	100562875-100562954,+	1.29	h	m r	m r	-	hsa-mir-323	ANMH46
HP-34	chr14	100563199-100563264,+	0.25	h	-	-	-	hsa-mir-323	
HP-35	chr14	100565724-100565804,+	1.40	h	m r	m r	r	hsa-mir-323	B209, X61
HP-36	chr14	100569845-100569926,+	1.67	-	m r	m r	r	hsa-mir-368	X124
HP-37	chr14	100576159-100576238,+	1.31	h	m r	m r	m r	hsa-mir-368	
HP-38	chr14	100576535-100576620,+	0.98	-	m r	m r	m r	hsa-mir-376a	ANMH47, X205
HP-39	chr14	100580290-100580371,+	0.93	-	-	-	-	hsa-mir-381	
HP-40	chr14	100582545-100582628,+	1.28	h	m r	m r	-	hsa-mir-381	B212, X199
HP-41	chr14	100583411-100583488,+	1.17	h	m r	m r	m	hsa-mir-381	B213, X177
HP-42	chr14	100584748-100584838,+	0.30	h	m r	m r	m r	hsa-mir-381	B214
HP-43	chr14	100585651-100585730,+	1.10	-	-	-	-	hsa-mir-381	
HP-44	chr14	100588536-100588615,+	1.11	h	m r	m r	m	hsa-mir-382	
HP-45	chr14	100591505-100591585,+	1.41	m	m r	m r	m	hsa-mir-134	ANMH51, B217, X23
HP-46	chr14	100596673-100596749,+	1.34	-	m r	m r	m r	hsa-mir-154	ANMH53, B219, X161
HP-47	chr14	100601390-100601468,+	0.74	h m	m r	m r	-	hsa-mir-369	ANMH55, B221
HP-48	chr14	100601548-100601614,+	0.18	m	m r	m r	-	hsa-mir-369	ANMH56
HP-49	chr14	100602002-100602081,+	0.78	h m	m r	m r	m r	hsa-mir-369	ANMH57, B565, X26
HP-50	chr15	29148489-29148618,-	1.15	-	m	-	-	hsa-mir-211	
HP-51	chr15	61945887-61945958,-	0.69	-	-	-	-	hsa-mir-422a	
HP-52	chr15	61956585-61956672,-	1.23	-	-	-	-	hsa-mir-422a	
HP-53	chr15	77283739-77283834,+	2.79	-	-	-	-	hsa-mir-184	
HP-54	chr15	77286586-77286662,+	0.19	-	-	-	-	hsa-mir-184	
HP-55	chr15	86952348-86952421,+	0.43	-	r	r	-	hsa-mir-7-2	
HP-56	chr16	14305325-14305407,+	1.25	h	m	m	m	hsa-mir-365-1	B707
HP-57	chr16	55455432-55455558,+	8.08	-	m	m	-	hsa-mir-138-2	
HP-58	chr16	65802725-65802850,-	1.03	-	r	-	-	hsa-mir-328	
HP-59	chr17	1557905-1558005,-	2.40	-	r	r	-	hsa-mir-22	
HP-60	chr17	7077268-7077349,-	0.04	-	-	-	-	hsa-mir-324	
HP-61	chr17	24212514-24212583,-	0.83	h m	m r	m r	m r	hsa-mir-144	ANMH67, B919, X65

HP-62	chr17	26925895-26925998,+	0.15	-	-	-	-	hsa-mir-365-2	
HP-63	chr17	54574700-54574820,-	4.57	-	-	-	-	hsa-mir-301	
HP-64	chr17	54583611-54583698,-	1.96	-	m	-	-	hsa-mir-301	
HP-65	chr17	55276486-55276565,+	1.08	-	-	-	-	hsa-mir-21	
HP-66	chr17	76710147-76710253,-	0.16	-	-	-	-	hsa-mir-338	
HP-67	chr18	54262074-54262128,+	0.20	-	-	-	-	hsa-mir-122a	
HP-68	chr19	10794434-10794541,-	4.81	-	m r	m	-	hsa-mir-199a-1	
HP-69	chr19	50828807-50828889,-	1.45	-	-	-	-	hsa-mir-330	
HP-70	chr19	58987053-58987142,+	0.42	-	-	-	-	hsa-mir-373	
HP-71	chr19	58991703-58991827,+	4.12	-	m r	m r	m	hsa-mir-373	
HP-72	chr20	61288007-61288074,+	0.01	-	-	-	-	hsa-mir-124a-3	
HP-73	chr20	16826662-16826746,+	0.07	-	-	-	-	hsa-mir-99a	
HP-74	chr21	25875807-25875895,+	0.52	-	-	-	-	hsa-mir-155	
HP-75	chr22	18387174-18387296,+	5.67	-	m	m	-	hsa-mir-185	
HP-76	chrX	49466772-49466857,+	0.09	-	-	-	-	hsa-mir-188	
HP-77	chrX	49476605-49476676,+	0.32	h m	m r	m r	-	hsa-mir-188	B121, X144
HP-78	chrX	49477366-49477449,+	0.20	-	m r	m r	m	hsa-mir-188	B122
HP-79	chrX	65017860-65017962,+	0.81	-	-	-	-	hsa-mir-223	
HP-80	chrX	73289960-73290065,-	2.01	h	m r	-	-	hsa-mir-374	
HP-81	chrX	84968305-84968369,-	0.14	-	-	-	-	hsa-mir-361	
HP-82	chrX	113872348-113872431,+	0.28	-	m r	m	-	hsa-mir-448	
HP-83	chrX	122415494-122415578,-	0.39	-	-	-	-	hsa-mir-220	
HP-84	chrX	133019128-133019229,-	1.95	-	-	-	-	hsa-mir-92-2	
HP-85	chrX	133028921-133029009,-	1.14	h m	m r	m r	-	hsa-mir-92-2	ANMH3, B343, X211
HP-86	chrX	133400068-133400148,-	1.59	h	m r	m r	-	hsa-mir-450	X138
HP-87	chrX	133405878-133405948,-	0.80	h	m r	m r	-	hsa-mir-424	B973, X141
HP-88	chrX	150798672-150798748,-	1.02	h	m	m	-	hsa-mir-224	AMH1, X154
HP-89	chrX	151232472-151232559,-	2.56	-	m r	m	-	hsa-mir-105-2	