

Chr	Scaffold	Start	End	Size
9	Hmel209007	4344840	4364959	20119
15	Hmel215006	1190144	1196212	6068
18	Hmel218003	221730	429239	207509
21	Hmel221012	779541	796444	16903
2	Hmel202004	1537111	1543050	5939
21	Hmel221001	297653	480497	182844
2	Hmel202006	1791672	1816820	25148
2	Hmel202006	1804029	1835608	31579
2	Hmel202006	4165194	4171714	6520
15	Hmel215047	2849864	2859210	9346
19	Hmel219018	354635	361054	6419

BayeScan log(PO)	BayPass mean XtX	Freq in <i>H. melpomene</i>	Freq in <i>H. cydno</i>
1.7222	7.95239143	0	0.93
1.8414	8.78515118	0	1
1.894	8.75630075	0	1
1.72	8.35788884	0	0.93
Not Significant	3.97886586	0.65	0.71
Not Significant	3.56709747	0.55	0.5
1.5501	Not Significant	0	0.79
1.0762	Not Significant	0.05	1
0.99213	Not Significant	0	0.93
0.97681	Not Significant	0.05	1
1.2169	Not Significant	0	1

Hmel2 annotation

HMEL009267

intergenic (upstream of *cortex*)

OBP41

HMEL013558

HMEL013559

HMEL003174

HMEL003175

HMEL003862

HMEL003863

HMEL016617

HMEL016621

HMEL016620

intergenic

HMEL011045

HMEL011044

HMEL011042

HMEL011043

HMEL011041

HMEL011040

HMEL011037

HMEL011039

HMEL011038

HMEL015626

HMEL015625

HMEL015624

HMEL012374

HmGr58

HmGr59

HmGr60

HmGr61

HmGr64

HmGr65