

Pseudo-observed models

Model supports

	SC_AC	SC_ACBD	SC_BD	SI	Classification error
SC_AC	4920	32	1	47	0.016
SC_ACBD	51	4862	60	27	0.0276
SC_BD	9	27	4897	67	0.0206
SI	31	20	24	4925	0.015

SC_AC: Secondary contact only between population A and C.
SC_BD: Secondary contact only between population B and D.
SC_ACBD: Secondary contact between population A and C, and between B and D.
SI: no gene flow.