

Taxon Set	95% lower boundary (observed gene to gene distance)	80% critical value upper boundary	Difference	Implied <i>p</i> value
None removed	34	11	23	< 0.05
14 Hybrids* BDF FRS HLM ICH IND MEX NGM RBD WGL AMB KLP PAL NIP MIC	5	5	<b>0</b>	<b>&gt; 0.05</b>
Random1: 17 21 14 13 26 27 18 1 7 4 5 24 19 9	12	4	8	< 0.05
Random2: 13 27 19 16 22 21 20 23 26 17 25 24 9 12	10	4	6	< 0.05
Random3: 24 22 15 23 12 17 26 8 20 18 12 3 27 10	18	4	14	< 0.05
Random4: 8 24 15 22 23 13 18 19 21 3 5 17 25 4	17	4	13	< 0.05
Random5: 12 2 18 6 15 3 9 17 20 4 16 7 27 24	13	4	9	< 0.05
Random6: 3 22 1 7 18 6 19 10 20 15 11 13 12 5	12	4	8	< 0.05
Random7: 15 4 27 12 2 23 13 6 14 21 20 18 17 26	10	3	7	< 0.05
Random8: 7 12 21 6 22 5 15 9 10 25 14 19 23 4	13	4	9	< 0.05
Random9: 20 4 21 18 15 17 1 14 6 22 16 9 5 8	10	4	6	< 0.05
Random10: 23 26 17 20 15 7 8 18 14 21 13 22 6 11	11	4	7	< 0.05