

Table S1. Reference DNA sequences selected for the study of CMM.

CMM	Plant species	Sequence source	Major ITS2 haplotype	Amplicon (bp)	Variation sites	Reference
Herba Potentillae Chinensis	<i>Potentilla chinensis</i> Ser.	17 CMM collected from 6 locality, 2 plant specimens from 2 locality, 1 NIFDC reference material, 3 counterchecking specimens	CACGTCGTTGCCCTCCCAACCC CTTCGGGAGTTGGCTGGGACGG ATGATGGCCTCCCGTGTGCTCCG CCGCGCGGTTGGCATAAATAAC GAGTCATCGGCGGCAACGCCG CGACAATCGGTGGTTGTCAAAC CTCGGTGTCCTGTGCGCGCGC GTCGTCTCGGCTCTTCCGATCT AATGCGCGTCGGTTCGCCGGCG CTTTCAACG	212	22 sites: T26C, T62C, T184C, T195C, A49G, A182G, C64G, C165G, C66T, C69T, C152T, C163T, C171T, C175T, C198T, G91A, G177A, G201A, A96C, C156A, T164G, G193M, in/del 169, 174	[1]
Herba Potentillae Discoloris	<i>Potentilla discolor</i> Bunge	7 CMM from 2 locality, 1 plant specimen, 1 NIFDC reference	CACGTCGTTGCCCTCCCAACCC CTCCGGGAGTTGGGTGGGACGG ATGATGGCCTCCCGTGCCTCCG TCGCGCGGTTGGCATAAATAACA AGTCCTCGGCGGCAACGCCGC GACAATCGGTGGTTGTCAAACC	210	3 sites: C37G, G93T, T163C	[1]

		material, 2 counterchecking specimens	TCGGTGTCTGTGCGGTGCGAG TCGTCCGGGGCTTTTCCAATCTG ATGCGCGTCGATTCGTGCGGCGCT TTCAACG			
-	<i>Potentilla kleiniana</i> Wight et Arn.	Plant specimens	CACGTCGTTGCCCTCCCAACCC CTTCGGGAGTTGGCTGGGACGG ATGGTGGCCTCCCGTGTGCTCCG TCGCGCGGTTGGCATAAATAACA AGTCTCGGCGGCCAACGCCGC GACAATCGGTGGTTGTCAAACC TCGGTGTCTGTGCGCGCGCGG TCGTCCGGGGCTTTTTTTCAATC TAACGCGCGTCGCTCCGTCGAC GCTTTCAACG	212	-	[1]
Radix Pulsatillae	<i>Pulsatilla chinensis</i> (Bge.) Regel	13 CMM collected from 4 locality, 14 plant specimen collected from 3 locality, 1 NIFDC reference	CACACAGCGTCGCCCCACCAA AGCATTTGGATGGGGGCGGAAA TTGGCCCCCGAGCCCCCGGG CACGGTCGGCACAAATGTTGGC CCTCGGCGGCGAGCGTCGCGGT CAGCGGTGGTTGTA CTCTCATCC TCCAAAGACAAAATGACGCGTC CGCCTCGTCGCCCCGCTGGGCGA	219	8 sites: C17T, C61T, C63T, C112T, C126G, G170A, G193T, A201T, in/del 64	[1,2]

		material, 2 counterchecki ng specimens, Accession no. GU732650	AGATGACCCAAGGAGTCTCCCC AACCGGAGACTTCCACCTG			
Radix Pulsatillae Cernuae	<i>Pulsatilla cernua</i> (Thunb.) Bercht.et Opiz.	Accession JN811070 & HQ829820	ACAGCGTCGCCCCACCAAAGC ATTTGGATGGGGGCGGAAATTG GCCCCCGAGCCCCCGGGGCA CGGTCCGCACAAATGTTGGCCC TCGGCGGCGAGCGTCGCGGTTA GCGGTGGTTGTACTCTCATCCTC CAAAGACAAAATGACGCGTCCG CCTCGTCGCCCCGCTGGGCAAAG ATGACCCAAGGAGTCTCCCCTA CCGGAGACTTCCACCTG	216	-	[1]
-	<i>Pulsatilla turczanino vii</i> Kryl. Et Serg.	Accession GU732649	CACACAGCGTCGCCCCACCAA AGCATTGGATGAGGGCGGAAA TTGGCCCCCGAGCCCTCCGGG GCACGGTCGGCACAAATGTTGG CCCTCGGCGGCGAGCGTCGCGG TCAGCGGTGGTTGTACTCTCATC CTCCAAAGACAAAATGACGCGT	219	-	[1]

			CCGCCTCGTCGCCCCGCTGGGCG AAGATGACCCAAGGAGTCTCCC CTACCGGAGACTTCCACCTG			
Radix Arnebiae	<i>Arnebia euchroma</i> (Royle) Johnst	11 CMM collected from 6 locality, 1 plant specimen, 1 NIFDC reference material, 1 counterchecki ng specimen, Accession no. EF199848, EF199860	CACATCGCGTCACCCCATCCAAA ATAATGTTGGATGTGGTGGATTG TGACCTCCTGTGTCTTGAGATGC AGTTGGTCGAAATTCGAGTCCG GAGCTTAGGACTTCACGACAAG TGGTGGTTGGATAACAACCTCGC GTCATGTCGTGTGCCAAGCCTCC GTGTCTCCGTAGACCCTAAGGC GCGTGCTTTCCAACCTCGTTCGTT GGGAAACCGTGCTACGACCG	223	5 sites: A27C, A109G, C143T, G153A, T189C	[1]
Radix Arnebiae	<i>Arnebia guttata</i> Bunge	Accession JX976805	CGCATCGCGTCACCCCATCCCAT GTAACATGGATGTGGTGGATTG TGACCTCCTGTGTCTTGAGATGC AGTTGGTCAAAATTTGAGTCTG GAGCTGAGGACTTCACGTCAAG TGGTGGTTGGATAACAACCTCGC	221	-	[1,3]

			GTCCTGTCGTGTGCCAAACCTCC ATTCCTCTGTGGACCCTAAGGCG TGTTTTTTTCCAACCTTGTGT GGGAAATCTTGCTACGA			
Radix Lithosper mi	<i>Lithosper mum erythrorhiz on Sieb. Et Zucc</i>	Plant specimens	CGCATCGCGTCACCCATCCCCAA CTCTCGTGGATGTGGTGGATTGT GACCTCCTGTGCCTTGCGGTGC AGTTGGTCGAAATTCGAGTTCG GAGTTAAGGACTTCACGACAAG TGGTGGTTGGACGACCAACTCG CGTCATGTCGTGTGCCAAACCTT CGTGGCTCTCAAGACCCTAAGG CACGTGCTACCCGACTCTTGGGT CGGAAAACCTCGTGCTACGACCG	224	-	[1]

References

1. Chen, S.L. *Standard DNA barcodes of Chinese Materia Medica in Chinese Pharmacopoeia*, 1st ed.; China Science Publishing & Media Ltd.: Beijing, China, 2015; pp. 168, 287, 446, 492.
2. Xie, L.; Wen, J.; Li, L.Q. Phylogenetic analyses of Clematis (Ranunculaceae) based on sequences of nuclear ribosomal ITS and three plastid regions. *Syst. Bot.* **2011**, *36*, 907–921.
3. Huang, J.F.; Zhang, M.L.; Cohen, J.I. Phylogenetic analysis of *Lappula Moench* (Boraginaceae) based on molecular and morphological data. *Plant. Syst. Evol.* **2013**, *299*, 913–926.