















































- Vassiljeva AN (1969) Critical notes on the genus *Parrya* R. Br. Notul Syst Herb Inst Bot Acad Sci KazSSR 6:27–31
- Vassiljeva AN (1974) Commentationes de systematica generis *Neurolooma* Andr. Notul Syst Herb Inst Bot Acad Sci KazSSR 8:27–31
- Warwick SI, Al-Shehbaz IA (2006) Brassicaceae: chromosome number index and database on CD-ROM. Pl Syst Evol 259:237–248
- Warwick SI, Sauder C (2005) Phylogeny of tribe Brassiceae (Brassicaceae) based on chloroplast restriction site polymorphisms and nuclear ribosomal internal transcribed spacer and chloroplast *trnL* intron sequences. Canad J Bot 83:467–483
- Warwick SI, Al-Shehbaz IA, Sauder C, Harris JG, Koch M (2004a) Phylogeny of *Braya* and *Neotorularia* (Brassicaceae) based on nuclear ribosomal internal transcribed spacer and chloroplast *trnL* intron sequences. Canad J Bot 82:376–392
- Warwick SI, Al-Shehbaz IA, Sauder C, Murray DF, Mummenhoff K (2004b) Phylogeny of *Smelowskia* and related genera (Brassicaceae) based on nuclear ITS DNA and chloroplast *trnL* intron sequences. Ann Mo Bot Gard 91:99–123
- Warwick SI, Francis A, Al-Shehbaz IA (2006a) Brassicaceae: checklist and database on CD-ROM. Pl Syst Evol 259:249–258
- Warwick SI, Sauder CA, Al-Shehbaz IA (2006b) Molecular phylogeny, morphology and cytological diversity of *Sisymbrium* (Brassicaceae). In: Sharma AK, Sharma A (eds) Plant genome: biodiversity and evolution: phanerogams (Angiosperm–Dicotyledons), vol 1C. Oxford and IBH, New Delhi with Science, USA, pp 219–250
- Warwick SI, Sauder CA, Al-Shehbaz IA, Jacquemoud F (2007) Phylogenetic relationships in the tribes Anchonieae, Chorisporae, Euclidieae, and Hesperideae (Brassicaceae) based on nuclear ribosomal ITS DNA sequences. Ann Mo Bot Gard 94:56–78
- Warwick SI, Mummenhoff K, Sauder CA, Koch MA, Al-Shehbaz IA (2010) Closing the gaps: phylogenetic relationships in the Brassicaceae based on DNA sequence data of nuclear ribosomal ITS region. Pl Syst Evol 285:209–232
- White TJ, Bruns T, Lee S, Taylor J (1990) Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ (eds) PCR protocols. Academic, New York, pp 315–322
- Yue JP, Sun H, Al-Shehbaz IA, Li JH (2006) Support for an expanded *Solms-laubachia* (Brassicaceae): evidence from sequences of chloroplast and nuclear genes. Ann Mo Bot Gard 93:402–411
- Yue JP, Sun H, Li J-H, Al-Shehbaz IA (2008) A synopsis of an expanded *Solms-laubachia* (Brassicaceae), and the description of four new species from western China. Ann Mo Bot Gard 95:520–538
- Yue JP, Sun H, Baum DA, Li JH, Al-Shehbaz IA, Ree R (2009) Molecular phylogeny of *Solms-laubachia* (Brassicaceae) s.l., based on multiple nuclear and plastid DNA sequences, and its biogeographic implications. J Syst Evol 47(5):204–411
- Zhao B, Liu L, Tan D, Wang J (2010) Analysis of phylogenetic relationships of Brassicaceae species based *Chs* sequences. Biochem Syst Ecol 38:731–739
- Zhou TY, Lu LL, Yang G, Al-Shehbaz IA (2001) Brassicaceae (Cruciferae). In: Wu ZY, Raven PH (eds) Flora of China 8, Science Press, Beijing, and Missouri Botanical Garden Press, St Louis, 1–193