

**Supplementary Table 2. The sequence conservation among Brp-MIR398a, Brp-MIR398b and Brp-CSD to its *Arabidopsis* homologs at amino acid level.**

miR398 and targets in <i>Arabidopsis</i>	Homologous in <i>Brassica rapa</i>	homology
ath-miR398a	Bra-MIR398a1	70.8%
	Bra-MIR398a2	74.7%
ath-miR398b	Bra-MIR398b1	64.4%
	Bra-MIR398b2	75.5%
ath-miR398c		
AtCSD1 (At1g08830)	BraCSD1-1 (Bra031642)	93.4%
	BraCSD1-2 (Bra018596)	34.7%
AtCSD2 (At2g28190)	BraCSD2-1 (Bra034394)	85.2%
	BraCSD2-2 (Bra011971)	61.1%
AtCCS (At1g12520)	BraCCS-1 (Bra016768)	87.5%
	BraCCS-2 (Bra026968)	53.1%