Ultra-dense SNP genetic map construction and identification of SiDt gene controlling the

determinate growth habit in Sesamum indicum L.

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**Supplementary Fig. S1.** QDt1 location for determinate growth habit on LG8 of the ultra-dense SNP map.

**Supplementary Fig. S2.** Amplification and screening of *Si*Dt 27-1 in F<sub>2</sub> population of the cross between Yuzhi DS899 and Ningbohei. M: DL 2,000 marker with the partial DNA bands of 250 bp and 100 bp; Lane 1-20: 92 bp amplicon of *Si*Dt 27-1 in *dt1* genotype individuals of F<sub>2</sub> population. Lane 21, 22, 28, 29, 31, 35 and 37: 97 bp amplicon of *Si*Dt 27-1 in *Dt* genotype individuals; Lane 23-27, 30, 32-34, 36, 38-40: amplicons of 92 bp and 97 bp in heterozygotic genotype (*Dt/dt1*).

**Supplementary Fig. S3**. Protein homology comparison of 3D structure between the SiDt and Sidt1 proteins. The SNP site of S79N is located at the junction region between different domains. Homology modelling shows that the 3D structures of SiDt (left) and Sidt1 (right) are same.

Supplementary Fig. S4. Phylogenetic analysis of *SiDt* homologues in sesame. Seven homologues of *Sis00046-1*, *Sis00076-1*, *Sis00121-1*, *Sis00157-1*, *Sis00157-2*, *Sis00154-1* and *Sis00141-1* are detected in the Yuzhi 11 genome (PRJNA315784) (version 2). Compared with SiDt, the protein identity of the 7 homologues ranges from 29%- 78%. The phylogenetic tree is constructed using MEGA 5.2 program according to the neighbor-joining (NJ) method. The line unit indicates the bootstrap value of 0.05.

Supplementary Fig. S5. Diversity analysis of *SiDt* among the sesame germplasm resources. Thirty core sesame germplasm are applied for the diversity analysis. A total of 18 SNP sites are detected in *SiDt* sequences among 5 accessions, i.e., M10, M11, M18, M19 and M29 as listed in Supplementary Table S10 online. Asterisks above the nucleotide sequences indicate the SNP sites of *SiDt* (*Sidt*) in Yuzhi 11, YuzhiDS899 and the 30 germplasm. The arrowhead indicates the SNP G397A site between Yuzhi 11 and Yuzhi DS899. The SNPs presented in more than 2 accessions are shaded in black. The

SNPs presented in one specific accession are shaded in grey. Bases in frame are exon sequences of SiDt.

**Supplementary Table S1.** Genome sequencing information of two parents and  $120 ext{ F}_2$  progeny in this study. Yuzhi DS899 as parent 1 is determinate (dt1 type), JS012 as parent 2 is indeterminate (the wild type). A total of  $120 ext{ F}_2$  progeny are randomly chosen from the  $ext{F}_2$  population of the cross between Yuzhi DS899 and JS012.

**Supplementary Table S2**. Genotyping matrix information of bin and SNP markers in the F<sub>2</sub> progeny. The information of 3,101 bins and 30,494 SNP markers are listed for linkage mapping.

**Supplementary Table S3.** Bin marker information of the ultra-density SNP genetic linkage map in sesame. 3,041 bin markers containing 30,193 SNP markers are located in the SNP linkage map.

Supplementary Table S4. Characteristics of the three phenotypes of determinacy growth habit in sesame. The determinacy trait of the five accessions are investigated at Sanya (109 50' E and 18 25' N), Pingyu (113 62' E and 32 97' N) and Yuanyang (113 97' E and 35 05' N) experimental stations in 2013. The average height trait data listed in the table are investigated at Yuanyang experimental station in 2013. Photoperiodic treatments of SD (12h light/12h dark) and LD (15h light/9h dark) are performed in growing chamber under the day/night temperature of 28  $^{\circ}$ C/24  $^{\circ}$ C with 70±1% relative humidity.

**Supplementary Table S5.** Location of QDt1 associated with the determinate growth habit in sesame using winQTLCart and QTLNetwork.

**Supplementary Table S6.** Annotation of all the genes in the QDt1 interval of Yuzhi DS899. Twenty five putative genes are detected in the physical distance corresponding to the QDt1 interval using BLASTP program. DS899s00170.026 and DS899s00170.027 listed in the table are adjacent genes to the QDt1 interval. \*'F' refers to forward direction and 'R' refers to reverse direction. KEGG refers to Kyoto Encyclopedia of Genes and Genomes.

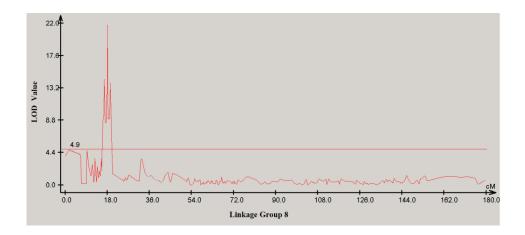
**Supplementary Table S7.** Genotypic variation and validation of the candidate SNP/InDel sites in sesame. Nine SNPs and 5 InDels are detected in the QDt1 interval. The *Si*Dt27-15 SNP and the *Si*Dt27-16 InDel listed in the table are detected at the adjacent flanking region of the QDt1 window. A total of 400 progeny of two F<sub>2</sub> populations of 'Yuzhi DS899 × JS012' and 'Yuzhi DS899 × Ningbohei' are used for screening target SNP/InDel marker.

**Supplementary Table S8.** Primer information of SiDt alleles for gene amplification and for real time PCR in sesame.  $\beta$ -Tubulin in sesame (SiTUB, JP631640) is used as the endogenous reference gene for qPCR.

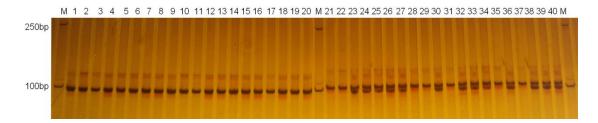
**Supplementary Table S9.** Global screening of SiDt homologues in the Yuzhi 11 genome. Seven homologues are detected using BLASTP, with the cutoff of E value  $\leq$ 1E-20 and the coverage of  $\geq$ 50%.

**Supplementary** Table S10. SiDt gene diversity analysis in sesame germplasm resources using PCR amplification. SiDt sequences in 30 sesame germplasm resources (M1-M30) are compared using DNAMAN software, in addition to the parents of the mapping population in the study.

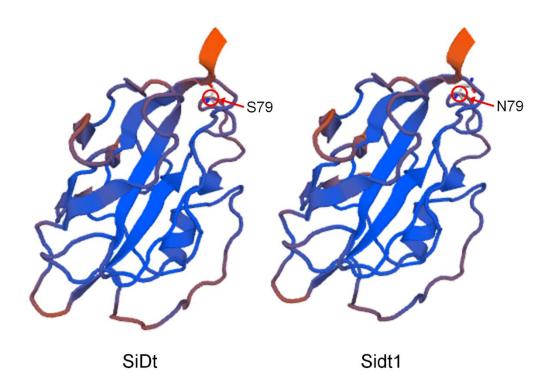
**Supplementary Table S11.** Diversity analysis of *SiDt* gene in sesame germplasm resources using public genome re-sequencing data. The public genome data of 715 accessionss are collected from PRJEB8078 in NCBI dataset.



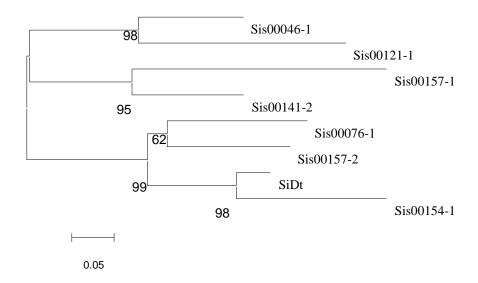
**Supplementary Fig. S1.** QDt1 location for determinate growth habit on LG8 of the ultra-dense SNP map. LOD refers to the Logistic Organ Dysfunction score. To determine the locus associated with the determinate trait locus, the LOD value is set at 4.9 using 1,000 permutations.



**Supplementary Fig. S2.** Amplification and screening of *Si*Dt 27-1 in F<sub>2</sub> population of the cross between Yuzhi DS899 and Ningbohei. M: DL 2,000 marker with the partial DNA bands of 250 bp and 100 bp; Lane 1-20: 92 bp amplicon of *Si*Dt 27-1 in *dt1* genotype individuals of F<sub>2</sub> population. Lane 21, 22, 28, 29, 31, 35 and 37: 97 bp amplicon of *Si*Dt 27-1 in *Dt* genotype individuals; Lane 23-27, 30, 32-34, 36, 38-40: amplicons of 92 bp and 97 bp in heterozygotic genotype (*Dt/dt1*).



**Supplementary Fig. S3**. Protein homology comparison of 3D structure between the SiDt and Sidt1 proteins. The SNP site of S79N (in red cycle) is located at the junction region between different domains. Homology modelling shows that the 3D structures of SiDt (left) and Sidt1 (right) are same.



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## Supplementary Fig. S5-1

	*	
Yuzhi 11	ATGGCAAAAATGTCATCGGACCCCTTGTGATCGGTAGGGTGGTCGGAGACGTTGTCGATCATTTCTCCTCCACCGTCAAAATGTCAGTCA	91
Yuzhi_Ds899	ATGGCAAAAATGTCATCGGACCCCCTTGTGATCGGTAGGGTGGTCGGAGACGTTGTCGATCATTTCTCCTCCACCGTCAAAATGTCAGTCA	91
M10	ATGGCAAAAATGTCATCGGACCCCCTTGTGATCGGTAGGGTGATCGGTTGTCGATCATTTCTCCTCCACCGTCAAAATGTCAGTCA	91
M11	ATGGCAAAAATGTCATCGGACCCCCTTGTGATCGGTAGGGTGATCGGAGACGTTGGTCGATCATTTCTCCTCCACCGTCAAAATGTCAGTCA	91
M18 M19	ATGGCAAAAATGTCATCGGACCCCCTTGTGATCGGTAGGGTGATCGGAGACGTTGTCGATCATTTCTCCTCCACCGTCAAAATGTCAGTCA	91 91
M29	ATGCCAAAATGTCATCGGACCCCCTTGTGATCGGTAGGGTGATCGGAGACGTTGTCGATCATTCTCCTCCACCGTCAAAATGTCAGTCA	91
114.5		71
Yuzhi 11	CTTACAACTCCAACAAGCATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA  CTTACAACTCCAACAAGACATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182 182
Yuzhi_Ds899 M10	CTTACAACTCCAACAAGCATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182
M11	CTTACAACTCCAACAAGCATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182
M18	CTTACAACTCCAACAAGGATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182
M19	CTTACAACTCCAACAAGCATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182
M29	CTTACAACTCCAACAAGCATGTCTACAATGGCCATGAGCTCTTCCCTTCCACAGTCACCTCTAAACCTAGGGTTGAGGTCCATGGTGGTGA	182
Yuzhi 11	TATGAGATCATTTTTCACCCTGCTAACATAATTTTTACTAATGATTTTTTTACACACCCCACTACTTCTTCTTCTAATTTCTGCTGATATT	273
Yuzhi_Ds899	TATGAGATCATTTTTCACCCTGCTAACAATAATTTTTACTAATGATTTTTTTACACACCCCCACTACTTCTTCTTCTAATTTCTGCTGATATT	273
M10	TATGAGATCATTTTTCACCCTGGTAACAATGATTTTTACTAATGATTTTTTACACACCCCCACTACTTCTTTCT	273
M11 M18	TATGAGATCATTTTCACCCTGTAACAATAATTTTACTAATGATTTTTTACACACCCCCACTACTTCTTCTTCTAATTTCTGCTGATATT TATGAGATCATTTTTCACCCTGTAACAATAATTTTACTAATGATTTTTTACACACCCCCACTACTTCTTCTAATTTCTGCTGATATT	273 273
M19	TATGAGATCATTTTCACCCTGETAACAATAATTTTACTAATGATTTTTTACACACCCCACTACTTCTTCTTCTTAATTTCTGCTGATATT	273
M29	TATGAGATCATTTTTCACCCTGTAACAATAATTTTACTAATGATTTTTTACACACCCCACTACTTCTTTCT	273
Yuzhi 11	$\tt TTTTTGGGAGTTTGTGTAAATTAAAGTAGTACGTGCAAATATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA$	364
Yuzhi_Ds899	$\tt TTTTTGGGAGTTTGTGTAAATTAAAGTAGTACGTGCAAATATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA$	364
M10 -	$\tt TTTTTGGGAGTTTGTGTAAATTAAAGTAGTACGTGCAAATATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA$	364
M11	TTTTTGGGAGTTTGTGTAAATTAAAGTAGTACGTGCAAATATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA	364
M18	TTTTTGGGAGTTTGTGTAAATTAAAGTAGTACGTGCAAATATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA	364
M19 M29	thm:thm:thm:thm:thm:thm:thm:thm:thm:thm:	364 364
F12.9	TITITOOOAGITTOTGTAAATTAAAGTAGTAGTAGTAGTATTTCGTGGGTTTTCAGTAATAATTTCCTCGTTTATATATA	304
Yuzhi 11	CATCATGACAGACCCTGATGTTCCTGGTCCTAGTGATCCATATCTGAGGGAGCACCTGCACTGCTATGCTTTCATTTTTAACTGCTTAAGA	455
Yuzhi_Ds899	GATCATGACAGACCCTGATGTTCCTGGTCCTAATGATCCATATCTGAGGGAGCACCTGCACTGGTATGCTTTCATTTTTAACTGCTTAAGA	455 455
M10 M11	GATCATGACAGACCCTGATGTTCCTGGTCCTAGTGATCCATATCTGAGGGAGCACCTGCACTGGTATGCTTTCATTTTTAACTGCTTAAGA GATCATGACAGACCCTGATGTTCCTGGTCCTAGTGATCCATATCTGAGGGAGCACCTGCACTGGTATGCTTTCATTTTTAACTGCTTAAGA	455
M18	GATCATGACAGACCCTGATGTTCCTGGTCCTAGTGATCCATATCTGAGGGACCACTGCACTGCTATGCTTTCATTTTTTAACTGCTTAAGA	455
M19	CATCATGACAGACCCTGATGTTCCTGGTCCTAGTGATCCATATCTGAGGGAGCACCTGCACTGGTATGCTTTCATTTTTAACTGCTTAAGA	455
M29	datcatgacagaccctgatgttcctggtcctagtgatccatatctgagggagcacctgcactggtatgctttcatttttaactgcttaaga	455
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Yuzhi 11	CCTGATTGATTTAATAAACTAGTATTCTTCAAAAAAACTAGAGTTTTGGAATGATCAACACCCTTTTCTCATCCAAGAGGGTAATCAACCAC	546
Yuzhi_Ds899	CCTGATTGATTTAATAAACTAGTATTCTTCAAAAAACTAGAGTTTTGGAATGATCAACACCCTTTTCTCATCCAAGAGGGGTAATCAACCAC	546
M10	CCTGATTGATTTAATAAACTAGTATTCTTCAAAAAACTAGGGTTTTTGGAAATGATCAACACCCCTTTTCTCATCCAAGAGGGTAATCAACCAC	546
M11 M18	CCTGATTGATTAATAAACTAGTATTCTTCAAAAAACTACAGTTTTGGAATGATCAACACCTTTTTCTCATCCAAGAGGGTAATCAACCACCCTGATTGAT	546 546
M19	CCTGATTGATTAATAAACTAGTATTCTTCAAAAAACTA AGTTTTGGAATGATCAACACCTTTTTCTCATCCAACAGGGTAATCAACACC	546
M29	${\tt CCTGATTGATTTAATAAACTAGTATTCTTCAAAAAAACTACAGTTTTTGGAATGATCAACACCTTTTTCTCATCCAAGAGGGTAATCAACCACCACCACCACCACCACCACCACCACCACCA$	546
Yuzhi 11	** AACACACTTTTTAGGGTTTTTTCTTTTAGTTTTTTTAAAAAATTATGGTAGAAACCGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA	637
Yuzhi_Ds899	$\texttt{AACACACTTTTTAGGGTTTTTTTTTTTTTTTTA} \underline{\texttt{A}} \texttt{AAAAATTATGGTAGAAACCGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA}$	637
M10	AACACACTTTTTAGGGTTTTTTCTTTTAGTTTTTTTTATAAAATTATGGTAGAAACCGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA	637
M11	AACACACTTTTTAGGGTTTTTTCTTTTAGTTTTTTTTTATAAAATTTATGGTAGAAACCGGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA	637 637
M18 M19	AACACACTTTTTAGGGTTTTTTTTTTAGTTTTTTTAT AAAATTATGGTAGAAACCGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA AACACACTTTTTAGGGTTTTTTTTTT	637
M29	AACACACTTTTTAGGGTTTTTTTCTTTTAGTTTTTTTA AAAATTATGGTAGAAACCGTTATTGAAATAGTCTATTGGAGGGGTCGATCATA	637
	-	
Yuzhi 11	${\tt TATATATATATATATGTATGTGTGTGGGTGGACTATAAAATTGAAGAGAAATTAATT$	728
Yuzhi_Ds899	${\tt TATATATATATATATGTATGTGTTGGGTGGACTATAAAATTGAAGAGAAATTAATT$	728
M10 -	${\tt TATATATATATATGTATGTGTTGGGTGGACTATAAAATTGAAGAGAAATTAATT$	728
M11	TATATATATATATATGTATGTGTTGGGTGGACTATAAAATTGAACAGGAAATTAATT	728
M18 M19	TATATATATATATATGTATGTGTTGGGTGGACTATAAAATTGAAGAGAAATTAATT	728 728
M29	TATATATATATATATGTATGTGTGTGGGCTGATAAAATTGAACAGAAATTAAATTTTAGAAAAAGGGAATTAAAACGTAATTCTAAATGC	728
	·	
Yuzhi 11	* AAACAAATTTAAATACTATTGCATTACAATTTTAAGATTTCTCTGCAAAAAACTAAGCTCCTAGGAAAGTTCGTTC	819
Yuzhi Ds899	ARACARATTTARATACTATTGCATTACAATTTTAAGATTTCTCTGCARAARACTRAGCTCCTAGGARAGTTCGTCACTTTAAGGTACAAG	819
M10	AAACAAATTTAAATTACTATTGCATTACAATTTTAAGATTTCTCCGCAAAAAACTAAGCTCCTAGGAAAGTTCTTCACCTTTAAGGTACAAG	819
M11	AAACAAATTTAAATACTATTGCATTACAATTTTAAGATTTCCCTGCAAAAAACTAAGCTCCTAGGAAAGTTCATTCA	819
M18	${\tt AAACAAATTTAAATACTATTGCATTACAATTTTAAGATTTCTCTGCAAAAAACTTAAGCTTCCTAGGAAAGTTCATTCA$	819
M19 M29	AAACAAATTTAAATACTATTGCATTACAATTTTAAGATTTCCTGCAAAAAACTAAGCTCCTAGGAAAGTTCATTCA	819 819
_ 340 0		0.20
Yuzhi 11	ATGTATTATAAAATTTTTTTGGTGATAATACATGAGATTCGTAAGTTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC	910
Yuzhi Ds899	ATGTATTATAAAATTTTTTTGGTGATAATACATGAGATTCGTAAGTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC	910
M10 -	$\tt ATGTATTATAAAATTTTTTTGGTGATAATACATGGGATTCGTAAGTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC$	910
M11	ATGRATTATAAAATTTTTTTGGTGATAATACATGAGATTGGTAAGGTTAGGGTGATGGTCTACACATGTGATGTAAGGTCCACAAATC	910
M18 M19	ATGTATTATAAAATTTTTTTGGTGATAATACATGAGATTCGTAAGTTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC ATGTATTATAAAATTTTTTTTGGTGATAATACATGAGATTCGTAAGTTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC	910 910
M29	ATGTATTATAAAATTTTTTTGGTGATAAACATGAGATTGGTAAGTTTAGAGTGATGGTCTACACTACATGTGATGTAAGGTCCACAAATC	910

## Supplementary Fig. S5-2

Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAAGTAGGTTTCATCGAGTCACTTACAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAAGTAGGTTTCATCGAGTCACTTACAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTCATCCTATGAAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC CCGCATCATATGTGTCAGACGCACTACGAAAAACCCTTGTTTCATCCTATGAAAAAGTAGGTTTCATCGAGTCACTTCCAACTAAATCTC	1001 1001 1001 1001 1001 1001
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	TGTCCAAGTGATGATCATGAAAATGTTGTTTGTCGTGTCGACTATCCTTCATTTCATTATCCTATGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTCGACTATCCTTCATTTCATTATTCCTATGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTCGACTATCCTTCATTTTCATTATCCTATTGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTGGACTATCCTTCATTTTTCATTATCCTATGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTCGACTATCCTTCATTTCATTATCCTTAGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTCGACTATCCTTCATTTCATTATCCTTATGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTGTCGTGTCGACTATCCTTCATTTCATTATCCTTATGGAAAATTACCCTTTAAAAATATA TGTCCAAGTGATGATCCATGAAAATGTTGTTTTTGTCGTGTCGACTATCCTTCATTTCATTATCCCTATGGAAAATTACCCTTTAAAAATATA	1092 1092 1092 1092 1092 1092 1092
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	*  CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGCAAATTACATAAAGCACA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGTGAAATTACATAAAGCACA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGT GAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGT GAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGT GAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGTTGAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGTTGAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGTTGAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTGACTGTTGAAATTACATAAAGCATA CTCTTGCCTGGTTCCACGTTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTTGACTGTTGAAATTACATAAAAGCATA CTCTTGCCTGGTTCCACGTTCTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTTGACTGTTGAAATTACATAAAAGCATA CTCTTGCCTGGTTCCACGTTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTTGACTGTTGAAATTACATAAAAGCATAA CTCTTGCCTGGTTCCACGTTTCTATTTCAACATCGTATTATCTGATTGTTGAATAGTGATAATTTTGACTGTTGAAATTACATAAAAGCATAA CTCTTGCCTGGTTCCACGTTTTTCAACATCGTATTATCTTGATTGTTGAATAGTGATAATTTTGACTGTTGAAATTACATAAAAACATAA CTCTTGCCTGGTTCCACGTTTTTCAATTTCAACATCGTATTATCTTGATTGTTGAATAGTGATAATTTTTTTT	1183 1183 1183 1183 1183 1183 1183
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	* ATATTAAAATAGAATAACTAAATCTGTATCGGATTCTCAATTGTTTACCCTAGTTAAAAAAATTTGCTTATATGTGGCTTGACAAATG ATATTAAAATAGAATAACTAAATCTGTATCGGATTCTCAATTGTTTACCCTAGTTAAAAAAAA	1274 1274 1274 1274 1274 1274 1274
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	* TACTTATAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACACTGACAAGTGCAGTTAACTAATGAATTACG TACTTATAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACACCTGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAACTAGGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG TACTT TAATTAATTAATCACATAAACATTTGACAAACAATGAGCCCTCTTTCAAAAACAC TGACAAGTGCAGTTAACTAATGAATTACG	1365 1365 1365 1365 1365 1365 1365
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	* TTGTTTTTATTTTACATATATGTACTAAATCTACTGACAAACATAAGTTGATTGA	1456 1456 1456 1456 1456 1456 1456
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	CCACAGATTCCTCATTCGGTATGAATAAAGATCATATATACTCCATTTAATTAA	1547 1547 1547 1547 1547 1547 1547
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	AATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGATTTCTTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGTACTACGAGGATGCCGAGGCCAAACATAGGAATACACAGGAATTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTAGGATAGGAGTACGAGGTCCGAGGCCAAACATAGGAATACACAGGAATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTTCTTGTTTACATGTAATTATCAATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAAATTTTTCTTGTTTACATGTACTTACATGTACTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAAATTTTTCTTGTTTACATGTACTTGCAGGAATACACAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAATTTTTCTTGTTTACATGTACTTCTGCAGGAAAAGAAGTAGTGAGCTACGAGATGCCGAGGCCAAACATAGGAATACACAGGAAATACACAGGAATTTTTTCTTGTTTACATGTACTTACATGTACTGCAGGAAAAGAAGTAGTAGGAATACACAGGAATACACAGGAATACACAGGAATTTTTCTTGTTTACATGTACTTACATGTACTGCAGAAAAGAAGTAGTAGGAATACACAGGAATACACAGGAAAGAAGAAGTAGTAGGAATGCAGAGGATGCCGAGGCCAAAAAGAAGAAGTAGGAATACACAGGAAAGAAGAAGAAGAAGAAGAAGTAGGAATGCAAAGGAATACACAGGAATACACAGGAATACACAGGAATACACAGAAAGAA	1638 1638 1638 1638 1638 1638
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	* TTTGCATTTGTGCTGTTCAAGCAGAAGAAGAGACAATTAGGATGCGTGAGGGCACCAGCTTGTAGGGATGGAT	1729 1729 1729 1729 1729 1729 1729
Yuzhi 11 Yuzhi_Ds899 M10 M11 M18 M19 M29	CCCAGGAAAATGAGTTAGGTCTCCCTGTTGCTGCTGTTTACTTCAATTGCCAGAGAGAG	1809 1809 1809 1809 1809 1809

Supplementary Fig. S5. Diversity analysis of SiDt among the sesame germplasm resources. Thirty

core sesame germplasm are applied for the diversity analysis. A total of 18 SNP sites are detected in *SiDt* sequences among 5 accessions, i.e., M10, M11, M18, M19 and M29 as listed in Supplementary Table S10 online. Asterisks above the nucleotide sequences indicate the SNP sites of *SiDt* (*Sidt*) in Yuzhi 11, YuzhiDS899 and the 30 germplasm. The arrowhead indicates the SNP G397A site between Yuzhi 11 and Yuzhi DS899. The SNPs presented in more than 2 accessions are shaded in black. The SNPs presented in one specific accession are shaded in grey. Bases in frame are exon sequences of *SiDt*.