

Supplemental Table S1 The gene-specific primes for selected DEGs

Gene ID	Primer	Sequence(5'–3')
101753893	Seita.3G110700-10G-qRTF	GGATGAAGCTCCTGCTCAAG
	Seita.3G110700-10G-qRTR	CGATGTTGCCGACGATGA
101754330	Seita.9G475300-11G-qRTF	GCCTGGAGCACATTAGCA
	Seita.9G475300-11G-qRTR	CCGTCGTAGAAGAACCTGAG
101757408	Seita.8G130200-28G-qRTF	GGTGGTGGACGATACCTTCA
	Seita.8G130200-28G-qRTR	CAGTGGTGTGCTTCCCTTT
101761991	Seita.3G143100-22G-qRTF	CGCTCGCCTACTATGGTGA
	Seita.3G143100-22G-qRTR	CACACGCCCTGACTCAATC
101764145	Seita.3G014700-4G-qRTF	GCCGTTCCAGAAGTTCATCA
	Seita.3G014700-4G-qRTR	TCCTTGTTCCAGCTCCAGTA
101767747	Seita.5G396500-35G-qRTF	CCTCCTCATCTCCGTCATTG
	Seita.5G396500-35G-qRTR	GAGCAGCCAGTAGAAGTAGTC
101771644	Seita.5G238500-32G-qRTF	CGGTGTACGACAAGGAGAC
	Seita.5G238500-32G-qRTR	CACCTGTATGCGGGGAAG
101772106	Seita.3G034900-8G-qRTF	GGACTCCACCTTAGAAGATGACT
	Seita.3G034900-8G-qRTR	CTGCTGTTCCCACCGAATATC
101772473	Seita.6G159400-9G-qRTF	ACACGCTGGACCAGATCA
	Seita.6G159400-9G-qRTR	CGTTGACGCAGAAGTAGAGC
101773925	Seita.9G440900-2G-qRTF	CAGGCAGAAGGAGCAGTTC
	Seita.9G440900-2G-qRTR	TGTTGACAGCGTGTGTGT
101774390	Seita.7G093500-14G-qRTF	CACCATCGGGAAGCAGAG
	Seita.7G093500-14G-qRTR	GTAGGCTCCGCACTGTAG
101775495	Seita.8G170100-13G-qRTF	AAGGTGACGAACCGTGCTA
	Seita.8G170100-13G-qRTR	CCGTTGGTGTGATCTTCTTG
101776721	Seita.8G200200-12G-qRTF	TGGCTGGAGCATGGAGTA
	Seita.8G200200-12G-qRTR	TGCTTGAACAGGACGAGTG
101779143	Seita.2G289500-20G-qRTF	CGGACGGCAAGTCATTCAT
	Seita.2G289500-20G-qRTR	GAGTTGAGCATCCAGGTGTAG
101784607	Seita.1G031400-33G-qRTF	GTGTTGAGACCAAGGTAGC
	Seita.1G031400-33G-qRTR	TGATGTCCTCGTCCCTAACC

Supplemental Table S2 Differentially expressed genes (DEGs) of two foxtail millet genotypes under control and low nitrogen conditions.

Part	Comparison	DEGs	Upregulated	Downregulated
Shoot	JG20LN vs JG20CK	4256	2214	2042
	JG22LN vs JG22CK	1291	431	860
	JG22CK vs JG20CK	3581	1346	2235
	JG22LN vs JG20LN	1132	457	675
Root	JG20LN vs JG20CK	3191	1862	1329
	JG22LN vs JG22CK	1881	1115	766
	JG22CK vs JG20CK	3852	1547	2305
	JG22LN vs JG20LN	2028	932	1096

Supplemental Table S3 Number and percentage of differentially expressed TFs between two different foxtail millet varieties in response to LN condition.

TF terms	Shoot JG20LN vs JG20CK	Shoot Common	Shoot JG22LN vs JG22CK	Root JG20LN vs JG20CK	Root common	Root JG22LN vs JG22CK
bHLH	18(12.5%)	3(16.7%)	6(16.2%)	8(7.1%)	1(3.0%)	6(5.7%)
bZIP	5(3.5%)	1(5.6%)	3(8.1%)	7(6.2%)	2(6.1%)	3(2.9%)
DIVARICATA	2(1.4%)	1(5.6%)	1(2.7%)	2(1.8%)	1(3.0%)	2(1.9%)
ERF	25(17.4%)	1(5.6%)	3(8.1%)	22(19.5%)	7(21.2%)	23(21.9%)
GATA	4(2.8%)	0(0.0%)	0(0.0%)	3(2.7%)	1(3.0%)	2(1.9%)
HHO	2(1.4%)	1(5.6%)	1(2.7%)	1(0.9%)	1(3.0%)	2(1.9%)
Hsf	5(3.5%)	1(5.6%)	3(8.1%)	2(1.8%)	1(3.0%)	3(2.9%)
IBH(S)/HBP(R)	2(1.4%)	1(5.6%)	1(2.7%)	1(0.9%)	1(3.0%)	1(1.0%)
MADS	4(2.8%)	2(11.1%)	4(10.8%)	4(3.5%)	0(0.0%)	2(1.9%)
MYB	12(8.3%)	2(11.1%)	6(16.2%)	6(5.3%)	1(3.0%)	7(6.7%)
MYC	1(0.7%)	0(0.0%)	0(0.0%)	0(0.0%)	0(0.0%)	1(1.0%)
NAC	1(0.7%)	1(5.6%)	1(2.7%)	0(0.0%)	0(0.0%)	1(1.0%)
NF-Y	6(4.2%)	0(0.0%)	0(0.0%)	4(3.5%)	3(9.1%)	4(3.8%)
PCF	3(2.1%)	0(0.0%)	0(0.0%)	1(0.9%)	1(3.0%)	2(1.9%)
RAX	2(1.4%)	0(0.0%)	0(0.0%)	0(0.0%)	0(0.0%)	2(1.9%)
TCP	2(1.4%)	0(0.0%)	0(0.0%)	1(0.9%)	0(0.0%)	0(0.0%)
TGAL	3(2.1%)	0(0.0%)	0(0.0%)	8(7.1%)	3(9.1%)	3(2.9%)
Trihelix	1(0.7%)	0(0.0%)	2(5.4%)	2(1.8%)	0(0.0%)	7(6.7%)
WRKY	29(20.1%)	0(0.0%)	0(0.0%)	11(9.7%)	6(18.2%)	25(23.8%)
others	17(11.8%)	4(22.2%)	6(16.2%)	30(26.5%)	4(12.1%)	9(8.6%)
Total	144(100%)	18(100%)	37(100%)	113(100.0%)	33(100.0%)	105(100.0%)

Supplemental Table S4 The DEGs related with Starch and sucrose metabolism in two varieties.

Parts	Shoots				Roots			
	LNJG20-CKJG20		LNJG22-CKJG22		LNJG20-CKJG20		LNJG22-CKJG22	
Treatments	UP	DOWN	UP	DOWN	UP	DOWN	UP	DOWN
Gene ID	101761744	101765726	101756806	101783222	101754475	101765726	101773502	101765726
Gene ID	101756806	101765746	101778895	101786399	101781847	101754626	101768458	101777643
Gene ID	101766064	101769822	101753890	101765726	101773502	101783222	101765746	105914669
Gene ID	101759340	101773215		101773215	101781445	101784113	101786613	101784144
Gene ID	101760963	101756027		101765746	101784519	101774056	101779998	101769336
Gene ID	101763600	101776290		101774746	101775037	101755255	101770526	101784175
Gene ID	101755745	101766477		101776688	101776290	101765110		101786459
Gene ID	101758779	101775884		101784175	101775239	101784144		101774056
Gene ID	101755613	101776688			101786844	101752753		
Gene ID	101786613	101754626			101772298	101759310		
Gene ID	101778895	101752753				101784175		
Gene ID	101781208	101785561				101774804		
Gene ID	101781159	101755626				101777786		
Gene ID	101782849	101768057				101754303		
Gene ID	101753874	101768040				101775414		
Gene ID	101781445	101769477				101786399		
Gene ID	101784519	101774746				101774746		
Gene ID	101753890	101768017				101768458		
Gene ID	101754475	101760405				101777643		
Gene ID	101764238	101758109						
Gene ID	105914669	101775414						
Gene ID		101782643						
Gene ID		101780809						
Gene ID		101786924						
Gene ID		101765386						

Supplemental Table S5 The DEGs related with photosynthesis in two varieties.

Gene ID	Gene Item	LNJG20-CKJG20	LNJG22-CKJG22
19526781	PsaA	DOWN	DOWN
101755060	PsaK	DOWN	DOWN
101759541	PsaN	DOWN	DOWN
101771677	PsaO	DOWN	DOWN
101779542	PsaG	DOWN	DOWN
101785215	PsaH	DOWN	DOWN
19526757	PsbC	DOWN	DOWN
101753687	PsbQ	UP	DOWN
101785134	PsbA	DOWN	DOWN
19526775	b	UP	DOWN
101767618	LHC	DOWN	DOWN

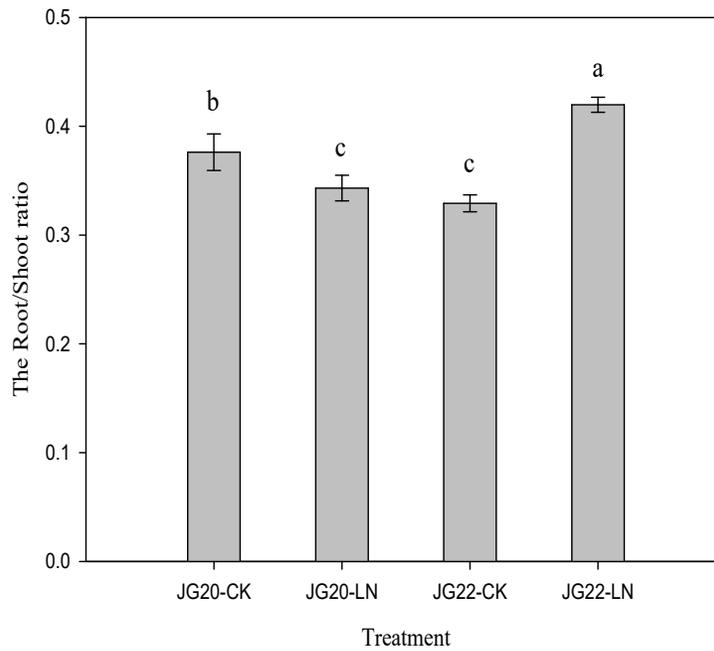
Supplemental Table S6 The DEGs related with Plant hormone signal transduction in two varieties.

gene_id	Gene Item	Shoot		Root		
		LNJG20-CKJG20	LNJG22-CKJG22	LNJG20-CKJG20	LNJG22-CKJG22	
101770000	AUX/IAA	DOWN	DOWN	DOWN	UP	
101763152	AUX/IAA	UP	DOWN	UP	UP	
101756773	AUX/IAA	UP	UP	UP	DOWN	
101786423	AUX/IAA	UP	UP	DOWN	UP	
101776449	AUX/IAA	UP	UP	UP	UP	
101776224	AUX/IAA	DOWN	UP	UP	UP	
101771394	AUX/IAA	DOWN	DOWN	DOWN	UP	
101776797	AUX/IAA	UP	UP	UP	UP	
101773098	AUX/IAA	DOWN	UP	UP	DOWN	
101782847	ARF	UP	UP	UP	UP	
101786550	ARF	UP	UP	UP	UP	
101759755	ARF	UP	UP	UP	UP	
101770279	Auxin	GH3	DOWN	DOWN	UP	UP
101786612		GH3	DOWN	DOWN	UP	UP
101777525		GH3	DOWN	DOWN	DOWN	UP
101768288		GH3	UP	UP	UP	UP
101768539		SAUR	DOWN	DOWN	UP	UP
101755080		SAUR	DOWN	DOWN	UP	UP
101757474		SAUR	DOWN	UP	UP	UP
101773025		SAUR	DOWN	UP	UP	UP
101786592	SAUR	DOWN	UP	DOWN	DOWN	
101759405	SAUR	UP	DOWN	UP	UP	
101774716	SAUR	DOWN	UP	UP	UP	
101776313	SAUR	DOWN	DOWN	DOWN	DOWN	
101776714	SAUR	DOWN	UP	DOWN	DOWN	
101784781	SAUR	UP	DOWN	UP	UP	
101768762	AHP	UP	DOWN	UP	UP	
101759627	AHP	UP	UP	UP	UP	
101773981	AHP	UP	UP	UP	UP	
101782248	Cytokinin	AHP	UP	UP	DOWN	DOWN
101785369		B-ARR	UP	UP	UP	UP
101781296		A-ARR	UP	UP	DOWN	UP
101767490		A-ARR	UP	DOWN	UP	UP
101778881	A-ARR	UP	UP	UP	UP	
101775649	Gibberellin	TF	DOWN	UP	UP	
101758377		EBF1/2	DOWN	DOWN	DOWN	DOWN
101778576	Ethylene	EIN3	UP	DOWN	DOWN	DOWN
101766570		EIN3	DOWN	DOWN	DOWN	DOWN
101785783	PYR/PYL	UP	UP	DOWN	UP	
101776342	Abscisic acid	PYR/PYL	UP	UP	UP	UP
101786686		PYR/PYL	UP	UP	DOWN	UP
101769449		PP2C	DOWN	DOWN	UP	DOWN
101753735	PP2C	DOWN	DOWN	UP	DOWN	

101783265	PP2C	DOWN	DOWN	DOWN	UP
101768564	PP2C	DOWN	DOWN	UP	DOWN
101780602	PP2C	DOWN	DOWN	DOWN	DOWN
101764618	SnRK2	DOWN	DOWN	DOWN	DOWN
101763654	SnRK2	DOWN	DOWN	DOWN	UP
101775125	SnRK2	DOWN	DOWN	DOWN	DOWN
101765115	SnRK2	UP	DOWN	UP	UP
101757593	ABF	DOWN	DOWN	UP	DOWN
101776070	ABF	DOWN	DOWN	DOWN	DOWN
101778442	ABF	DOWN	DOWN	UP	DOWN
101758999	ABF	UP	UP	UP	UP

Supplemental Table S7 The DEGs related with glycolysis in two varieties.

gene_id	Gene Item	Shoot		Root	
		LNJG20-CKJG20	LNJG22-CKJG22	LNJG20-CKJG20	LNJG22-CKJG22
101754626		DOWN	UP	DOWN	DOWN
101775414	H XK	DOWN	DOWN	DOWN	DOWN
101784144		DOWN	UP	DOWN	DOWN
101772215		DOWN	DOWN	DOWN	UP
101771409	A EP	UP	DOWN	UP	UP
101754303	G PI	UP	UP	DOWN	DOWN
101778700		DOWN	DOWN	DOWN	DOWN
101780397		DOWN	DOWN	DOWN	DOWN
101780059	A TP-PFK	DOWN	DOWN	UP	UP
101756322		UP	UP	DOWN	UP
101756917		UP	DOWN	DOWN	DOWN
101786189		UP	UP	UP	DOWN
101756517	F BA	DOWN	DOWN	DOWN	DOWN
101754112		DOWN	DOWN	DOWN	DOWN
101760718	T IM	UP	UP	UP	UP
101785151		DOWN	DOWN	DOWN	DOWN
101768835		DOWN	DOWN	DOWN	DOWN
101759360		DOWN	DOWN	DOWN	DOWN
101785907	P GK	UP	UP	UP	UP
101775268		DOWN	DOWN	DOWN	DOWN
101783820		DOWN	DOWN	UP	UP
101776203		DOWN	DOWN	DOWN	UP
101763575		UP	UP	UP	DOWN
101771740	P GMP	DOWN	DOWN	DOWN	DOWN
101757916		DOWN	UP	DOWN	DOWN
101781167		DOWN	DOWN	DOWN	UP
101769604	E NO	DOWN	DOWN	DOWN	DOWN
101783767	P CK	UP	UP	UP	DOWN
101768352		DOWN	DOWN	DOWN	DOWN
101759427		DOWN	DOWN	DOWN	DOWN
101769537	P K	UP	DOWN	UP	UP
101780488		DOWN	DOWN	DOWN	DOWN
101784708		UP	UP	UP	UP
101763108	P DH-E1 α	DOWN	DOWN	DOWN	UP
101775698	P DC	DOWN	DOWN	DOWN	DOWN
101763713		DOWN	DOWN	UP	UP
101769555	P DH-E2	UP	DOWN	UP	UP
101756978	D LD	DOWN	DOWN	UP	UP
101782764		DOWN	DOWN	UP	UP
101761749		UP	UP	UP	UP
101755454	A LDH	UP	UP	UP	DOWN
101776963		UP	UP	DOWN	UP
101762488		DOWN	UP	DOWN	UP



Supplemental Figure S1 The root/shoot ratio in two foxtail millet varieties under control and low nitrogen conditions. JG20-CK, JG20 under control; JG20-LN, JG20 under low nitrogen; JG22-CK, JG22 under control; JG22-LN, JG22 under low nitrogen. Error bars represented standard error of three biological replicates; Lowercase indicates significance at the level of $P < 0.05$.