

**Table1.** Juice quality analysis of sweet sorghum Genotypes of different origins

<b>Genotypes</b>	<b>FBY</b>	<b>SSWt</b>	<b>DSY</b>	<b>Brix</b>	<b>J.yld</b>	<b>Etoh</b>	<b>P</b>	<b>GR</b>	<b>SR</b>	<b>SC</b>	<b>TSS</b>
<b>Acc371st</b>	138.12	111.07	0.39	16.943	706	5763.95	82.72	23.32	3.125	13.4	16.2
<b>Acc402nw</b>	95.56	66.5	0.249	13.343	654	2532.98	85.56	20.29	3.125	15.4	18
<b>acc392st</b>	134.13	99.54	0.448	13.943	568	4130.69	88.16	15.3	2.05	13.4	15.2
<b>acc213st</b>	123.41	108.15	0.688	16.443	488	3698.67	84.35	29.4	4.166	14.17	16.8
<b>acc42sw</b>	96.23	78.24	0.463	15.043	398.45	2591.03	58.04	45.43	4.166	9.17	15.8
<b>acc1-1sw</b>	133.28	110.72	0.457	13.093	526.14	4261.16	85	16.34	2.5	15.3	18
<b>acc282st</b>	121.82	96.24	0.522	15.343	417.17	3554.99	87.86	25.41	3.125	12.3	14
<b>acc222st</b>	108.78	87.44	0.423	14.193	514.17	3391.03	76.9	24.19	3.125	12.92	16.8
<b>acc001et</b>	147.39	115.33	0.358	18.043	802.67	6474.19	79.88	18.41	2.5	13.56	17
<b>acc31sw</b>	119.95	96.43	0.419	16.543	447.54	4517.65	76	36.54	4.166	11.4	15
<b>acc002et</b>	127.85	101.72	0.437	15.343	441	4380.23	77.85	26.94	3.125	11.6	14.9
<b>acc421nw</b>	128.06	107.2	0.419	14.343	715.5	3985.67	70.37	32.89	3.125	9.5	13.5
<b>acc251st</b>	107.78	94.44	0.412	10.643	409.95	2541.31	59.06	39.19	4.166	10.63	18
<b>acc221st</b>	189.22	160.06	0.585	12.443	571.16	5542.17	68.5	22.81	2.5	10.96	16
<b>acc1331sw</b>	129.42	102.97	0.429	16.843	483.9	5167.69	67.33	41.25	4.166	10.1	15
<b>acc61sw</b>	119.1	97.61	0.419	15.843	460.29	4172.92	51.98	49.48	4.166	8.42	16.2
<b>acc264st</b>	103.67	83.56	0.523	19.043	463	3682.48	68.25	22.89	2.5	10.92	16
<b>acc301st</b>	166.28	138.95	0.528	15.643	601.38	6452.8	76.59	40.29	4.166	10.34	13.5
<b>acc201st</b>	88.52	71.72	0.312	16.293	679.17	3373.96	69.06	37.7	4.166	11.05	16
<b>acc412nw</b>	139.56	117.02	0.601	17.643	310.17	5113.04	77.5	31.5	3.125	9.92	12.8
<b>acc1-1sw</b>	164	133.95	0.512	15.593	607.48	5779.45	73.57	30.34	2.5	8.24	11.2
<b>acc71sw</b>	118.95	97.61	0.585	17.343	256.17	3770.74	62.4	33.39	3.125	9.36	15
<b>acc72sw</b>	118.89	91.53	0.436	16.043	705.17	3673.7	61.74	23.54	2.5	10.62	17.2
<b>acc32sw</b>	141.19	123.81	0.482	14.643	527.71	5499.78	67.29	27.32	3.125	11.44	17

<b>acc161st</b>	118.68	94.53	0.579	17.043	479.17	3677.81					
<b>acc41sw</b>	127.1	98.61	0.417	14.843	407.45	3962.43	54.3	29.14	2.5	8.58	15.8
<b>acc171st</b>	103.63	58.58	0.427	17.443	377	2114.81	68.8	40.37	4.166	10.32	15
<b>acc101nw</b>	106.48	94.67	0.422	15.943	545.87	4138.36	66.65	34.34	4.166	12.13	18.2
<b>acc381st</b>	160.11	127.98	0.489	15.243	560.12	5619.57	65.75	39.6	4.166	10.52	16
<b>acc351st</b>	129.06	102.17	0.574	16.843	513	4262.64	61.21	50.54	6.125	12.12	19.8
<b>acc141nw</b>	105.11	89.11	0.436	13.943	589	3155.15	100	40.6	4.166	10.26	10.26
<b>acc52sw</b>	92.22	70.28	0.515	18.693	424	2696.3					
<b>acc174st</b>	114.53	97.5	0.312	14.943	606	4508.49	57.87	48	4.166	8.68	15
<b>acc32sw</b>	138.57	118.07	0.427	14.043	667	4913.55	60.49	42.51	4.166	9.8	16.2
<b>acc18st</b>	132.33	160.78	0.545	18.043	715	7217.79	78.25	19.97	3.125	15.65	20
<b>acc91nw</b>	103.44	127.67	0.402	18.843	798	5812.03	69.33	33.76	4.166	12.34	17.8
<b>acc151nw</b>	106.41	122.61	0.313	15.393	803	5342.95	68.53	24.32	2.5	10.28	15
<b>acc312st</b>	74.34	89.06	0.471	16.243	360	2919.62	84.57	17.6	3.125	17.76	21
<b>acc342st</b>	102.72	122.41	0.558	12.393	402.67	2944.53	65.26	25.2	3.125	12.4	19
<b>acc31sw</b>	94.53	116.09	0.419	15.343	467.71	3920.73	69.16	38.86	4.166	10.72	15.5
<b>acc122nw</b>	81.61	105.9	0.376	13.643	373.67	3150.48	54.55	57.86	4.166	7.2	13.2
<b>acc362st</b>	108.33	134.5	0.544	14.393	557.67	4460.01	57.74	32.22	3.125	9.7	16.8
<b>acc391st</b>	94.33	114.45	0.406	16.893	383.35	4662.17	50.91	37.2	3.125	8.4	16.5
<b>acc241st</b>	86.34	107.5	0.391	16.243	427.12	3753.39					
<b>acc243st</b>	92.38	119.11	0.344	13.843	716.67	3572.05	61.19	43.28	3.125	7.22	11.8
<b>acct51sw</b>	92.72	122.81	0.673	11.843	658.67	2183.47	66.67	31.25	3.125	10	15
<b>Mn1383</b>	30.78	42.27	0.175	13.043	169.52	946.91	100	25.57	3.125	12.22	12.22
<b>Ramada</b>	37.11	49.77	0.203	13.443	190.96	1211.34	85	24.47	2.08	8.5	10
<b>Gkahon</b>	20.06	31.13	0.142	12.643	59.83	511.04					
<b>Gkgaba</b>	13.99	18.73	0.124	12.243	101.54	266.26	81.9	15.26	2.5	16.38	20

<b>Mn1500</b>	30.4	36.08	0.176	12.043	235.21	852.36	79.08	24.32	2.5	10.28	13
<b>Umbrella</b>	30.64	33.13	0.18	12.643	247.14	897.57	74.45	27.08	3.125	11.54	15.5
<b>Honey</b>	29.53	36.2	0.171	14.843	221.88	1040.81	76.47	36.32	4.166	11.47	15
<b>Rona</b>	95.61	119.56	0.248	13.443	475.67	4060.82	60.63	32.22	3.125	9.7	16
<b>EG002</b>	50.74	66.67	0.224	17.443	401.67	2472.39	46.04	36.93	4.166	11.28	24.5
<b>Brandes</b>	112.11	131.93	0.473	16.743	632.54	5327.99	58.65	48	4.166	8.68	14.8
<b>EG004</b>	91.28	123.33	0.409	16.443	468.32	4137.27					
<b>EG005</b>	45	52.89	0.23	15.443	268.81	1777.58					
<b>EG006</b>	66.43	88.58	0.329	19.043	353.79	3373.2	73.03	17.29	2.5	14.46	19.8
<b>Durado</b>	33.18	48.71	0.172	11.743	148.67	1064.35	62.49	29.63	4.166	14.06	22.5
<b>A22672</b>	50.78	53.44	0.248	12.843	356.21	1691.77					
<b>NGTJ-2</b>	62.44	74.11	0.293	17.043	430.04	2892.44					
<b>Karmifma</b>	104.11	129.11	0.448	16.043	611.13	4685.43					
<b>Acc006</b>	27.11	44.78	0.195	11.243	103.67	603.43					
<b>Melkam</b>	49.11	65.44	0.38	20.443	136.17	1965.9					
<b>acc003et</b>	38.78	50.44	0.239	18.043	114.17	1640.94					
<b>acc004et</b>	179.45	203.11	0.996	18.443	572.17	8329.75					
<b>acc005et</b>	161.45	194.45	0.661	21.243	479.17	10143.22					
<b>F-test</b>	<b>22.74**</b>	<b>123.25**</b>	<b>6.72*</b>	<b>22.35**</b>	<b>1245**</b>		<b>48.0**</b>	<b>28.14**</b>	<b>7.43*</b>	<b>31.3**</b>	<b>16.0**</b>
<b>LSD</b>	<b>8.25</b>	<b>56.20</b>	<b>0.09</b>	<b>7.10</b>	<b>968.50</b>	<b>964</b>	<b>14.03</b>	<b>10.33</b>	<b>1.78</b>	<b>3.66</b>	<b>8.03</b>
<b>CV</b>	<b>13.21</b>	<b>15.90</b>	<b>11.00</b>	<b>18.02</b>	<b>14.75</b>	<b>12.02</b>	<b>0.71</b>	<b>18.90</b>	<b>24.94</b>	<b>12.31</b>	<b>10.71</b>

\*\*Significant at 1% of probability ( $P < 0.01$ ); \* Significant at 5% of probability ( $0.05 = < p < 0.05$ ); ns - Not significant ( $P > 0.05$ ).  
 FBY: Fresh biomass yield; SSWt: Stripped stalk weight; DSY: Dry stalk yield; Brix: degree brix; J.yld: Juice yield; EtoH: Theoretical Ethanol yield; P:Purity%, GR: Glucose Recovery%; RS: Reducing sugar%; SC: Sucrose%; TSS: Total soluble sugar  
 L.S.D = Least Significant Difference. C.V = Coefficient of Variation.

**Table 3.** Estimates of genetic parameters of thirteen biofuel related quantitative traits in 68 sorghum accessions.

Traits	Grand	Range		SE	CV	PCV	GCV	H <sup>2</sup> (%)	GA	GAM%
	Mean	Min.	Max							
Purity %	68.37	46.04	100	3.37	2.13	0.33	0.252	61.05	25.09	36.70
Glucose Recovery (%)	33.54	15.26	57.86	0.75	13.75	0.33	0.252	51.94	10.81	32.23
Sugar recovery (%)	3.49	2.05	6.13	0.09	10.18	0.345	0.264	62.1493	2.14	61.32
Sucrose content (%)	10.83	7.2	17.76	0.17	17.42	1.02	0.792	77.2716	4.68	43.21
TotalSoluble Solids (%)	16.03	10	24.5	1.69	14.59	0.87	0.66	52.2801	5.57	34.75
Fresh biomass yield (Ton/ ha)	108.73	18.73	203.11	4.65	26.92	0.43	0.41	90.91	87.56	80.53
Stripped Stalk weight (Ton/ ha)	87.84	13.99	179.45	4.21	16.55	0.56	0.48	73.47	74.45	84.75
Dry Stalk yield (Kg/plant)	0.41	0.124	0.996	0	19.02	0.34	0.29	82.25	0.26	63.41
Brix (%)	15.43	10.64	21.24	0.18	6.48	0.21	0.2	50.7	6.05	39.21
Juice yield (gm/plant)	461.25	59.83	803	4.4	11.31	0.56	0.51	82.94	441.32	95.68
Ethanol yield (liter/ha)	3719.61	266.26	10143	18.92	22.75	0.54	0.52	61.13	2548.48	68.51

Note: SE= Standard error, CV= Coefficient of variance PCV = Phenotypic coefficient of variation, GCV = Genotypic coefficient of variation, H<sup>2</sup> = Broad sense heritability, GA= Genetic Advance, GAM= Genetic advance as percent of mean.