

Evaluation of improved lines in preliminary trials

Activity 1: yield trials in lentil.

Title	: Evaluating yield potential of lentil advanced lines in Marchouch and Kfardan												
Objectives	: Estimate the yield potential of lentil varieties and select lentil varieties for high performance and phenotypic stability.												
Activities	: Continued												
Expected outcomes	: High yield lines with beneficial traits including extra earliness, machine harvesting would be identified.												
Observations to be taken	: Plant height Height of the lowest pod Phenological traits Yield and yield components												
Trials	: Advanced Yield Trial- Large seeded (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial – Small seeded (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial- Early (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial- Machine harvest (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial- Micro Nutrient (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial – Extra early (33 genotypes + 2 improved checks + 1 local check) Advanced Yield Trial- Drought (33 genotypes + 2 improved checks + 1 local check)												
Key outcome	: Marchouch: The crop was not successful due to severe drought during early stage of the crop growth. However, seeds of individual genotypes were harvested and the trial will be repeated in the upcoming cropping season. Kfardan: There were significant genotypic differences in grain yield were found across all the trials conducted in Kfardan 2016. The best performing genotypes over the check variety are presented in Table 1,2,3,4,5, 6 and 7 as below. Table 1. The best performing genotypes over the check variety in AYT-LS2016 <table border="1" style="margin-left: auto; margin-right: auto;"> <thead> <tr> <th>Designation</th> <th>Pedigree</th> <th>Seed yield</th> </tr> </thead> <tbody> <tr> <td>x2011s17_20 _ 3</td> <td>BARIMASUR-6XLIRL-22-46-1-1-1-0</td> <td>1084</td> </tr> <tr> <td>x2011s139_124 _ 9</td> <td>FLIP97-29LXIG135395</td> <td>1038</td> </tr> <tr> <td>x2011s242_230 _ 3</td> <td>ILL10801XILL2711</td> <td>1008</td> </tr> </tbody> </table>	Designation	Pedigree	Seed yield	x2011s17_20 _ 3	BARIMASUR-6XLIRL-22-46-1-1-1-0	1084	x2011s139_124 _ 9	FLIP97-29LXIG135395	1038	x2011s242_230 _ 3	ILL10801XILL2711	1008
Designation	Pedigree	Seed yield											
x2011s17_20 _ 3	BARIMASUR-6XLIRL-22-46-1-1-1-0	1084											
x2011s139_124 _ 9	FLIP97-29LXIG135395	1038											
x2011s242_230 _ 3	ILL10801XILL2711	1008											

x2011s33_34_32	ILL10731XILL9889	1002
x2011s17_20_2	BARIMASUR-6XLIRL-22-46-1-1-1-0	965
x2011s126_116_21	ILL4400X28509	917
x2011s133_119_4	FLIP97-29LXFLIP97-33L	913
Check	ILL4400	688
Trial mean		790.30
LSD(<0.001 level)		181.78
Heritability		0.74

Table 2. The best performing genotypes over the check variety in AYT-SS2016

Designation	Pedigree	Seed yield
x2011s_246_25	ILL10749XILL3597	1099
x2011s_183_16	ILL7115XILL2585	1090
x2011s_203_2	ILL10749XILL3597	1073
x2011s_192_45	ILL10749XILL7979	1072
x2011s_176_1	ILL4605XILL6024	1062
Check	Check	733
Trial mean		805.57
LSD(<0.001 level)		287.51
Heritability		0.68

Table 3. The best performing genotypes over the check variety in AYT-E 2016

Designation	Pedigree	Seed yield
x2011s_129_13	FLIP96-49LXFLIP97-33L	1178
x2011s_119_25	FLIP97-29LXFLIP97-33L	1163
x2011s_206_26	ILL10800X33109	1141
x2011s_130_1	ILL4402XILL7979	1104
x2011s_129_36	FLIP96-49LXFLIP97-33L	1096
x2011s_129_28	FLIP96-49LXFLIP97-33L	1096
x2011s_122_26	ILL1712XILL10072	1048
x2011s_247_19	LIRL-22-46-1-1-1-1-0XAKM 302	993
x2011s_172_20	ILL7723XADA'A	978
x2011s_203_2	ILL10749XILL3597	970
x2011s_171_7	Barimusor- 6xL-7713	959
x2011s_138_20	Barimusor 6xILL7979	933
Check	ILL8006	720
Trial mean		893.90
LSD(<0.001 level)		208.08
Heritability		0.79

Table 4. The best performing genotypes over the check variety in AYT-EE2016

Designation	Pedigree	Seed yield
x2013_20_36	BARIMASUR-6XLIRL-22-46-1-1-1-0	1219
x2013_20_7	BARIMASUR-6XLIRL-22-46-1-1-1-0	1193
x2013_20_26	BARIMASUR-6XLIRL-22-46-1-1-1-0	1133
x2013_125_15	FLIP97-34LXFLIP97-33L	1126
x2013_126_54	FLIP97-34LXFLIP97-33L	1119
x2013_21_2	BARIMASUR-6XLIRL-22-46-1-1-1-0	1104
x2013_118_3	FLIP97-29LXFLIP97-33L	1081
x2013_20_3	BARIMASUR-6XLIRL-22-46-1-1-1-0	1052
x2013_175_35	ILL4605XILL6024	1037
x2013_72_28	ILL7978XILL7537	1032
x2013_125_40	FLIP97-34LXFLIP97-33L	1022
x2013_119_14	FLIP97-29LXFLIP97-33L	1007
x2013_82_10	KHAJURA-2XILL5888	1007
x2013_126_5	FLIP97-34LXFLIP97-33L	1007
x2013_126_8	FLIP97-34LXFLIP97-33L	988
x2013_119_24	FLIP97-29LXFLIP97-33L	963
x2013_280_18	Barimusor- 6XILL6994	956
x2013_72_19	ILL7978XILL7537	956
x2013_166_8	ILL4605XILL358	944
x2013_140_1	ILL4404XILL7950	933
x2013_266_2	LIRL-22-46-1-1-1-0XAKM 302	932
x2013_142_15	ILL4605XILL7950	926
Check	Check	681
Trial mean		945
LSD(<0.001 level)		216.24
Heritability		0.78

Table 5. The best performing genotypes over the check variety in AYT-MH 2016

Designation	Pedigree	Seed yield
x2011s_200_13	ILL10801XILL7950	1093
x2011s_176_1	ILL4605XILL6024	1073
x2011s_199_9	ILL10801XILL7979	1065
x2011s_243_12	ILL10750XILL6994	1024
x2011s_200_10	ILL10801XILL7950	1014
Local check	Check	722
Trial mean		834.01
LSD(<0.001 level)		224.08
Heritability		0.74

Table 6. The best performing genotypes over the check variety in AYT-DT2016

Designation	Pedigree	Seed yield
x2011s_119_26	FLIP97-29LXFLIP97-33L	1207
x2011s_172_34	ILL7723XADA'A	1067
x2011s_125_23	FLIP97-34LXFLIP97-33L	963
x2011s_206_58	ILL10800X33109	947
x2011s_223_6	ILL10800XILL7201	921
x2011s_195_4	ILL10750XILL7950	904
x2011s_111_26	ILL8007XILL759	904
x2011s_123_36	ILL1712XILL10072	889
x2011s_163_9	ILL4402XILL5588	881
x2011s_204_51	ILL10750X33108	875
x2011s_119_23	FLIP97-29LXFLIP97-33L	871
Local check	Check	637
Trial mean		802
LSD(<0.001 level)		235.05
Heritability		0.69

Table 7. The best performing genotypes over the check variety in AYT-MN 2016

Designation	Pedigree	Seed yield
x2011s_163_15	ILL4402XILL5588	1259
x2011s_126_60	FLIP97-34LXFLIP97-33L	1167
ILL6821	ILL6821	1139
x2011s_97_20	ILL5883XILL10750	1117
x2011s_125_36	FLIP97-34LXFLIP97-33L	1116
x2011s_198_33	ILL10801XILL7979	1094
x2011s_97_17	ILL5883XILL10750	1083
x2011s_192_45	ILL10749XILL7979	1066
x2011s_54_14	ILL8595XILL10741	1058
x2011s_118_21	FLIP97-29LXFLIP97-33L	1011
x2011s_278_1	ILL975XILL10657	1003
x2011s_89_25	ILL5883XILL8	979
x2011s_118_12	FLIP97-29LXFLIP97-33L	955
x2011s_176_31	ILL4605XILL6024	937
x2011s_110_13	ILL8007XILL759	904
x2011s_72_44	ILL7978XILL7537	904
x2011s_115_22	ILL975X28505	879
x2011s_175_1	ILL4605XILL6024	831
x2011s_172_25	ILL7723XADA'A	817
x2011s_63_7	ILL3796XILL4605	816
x2011s_82_3	KHAJURA-2XILL5888	793
x2011s_244_9	ILL10800XILL4637	786
Check	Check	539

		Trial mean		862.10
		LSD(<0.001 level)		242.14
		Heritability		0.81

Activity 2 : Preliminary yield trials in Faba bean

Title	:	Evaluating yield potential of faba bean advanced lines.																																				
Objectives	:	Estimate the yield potential of varieties, select faba bean varieties for high performance and forward to advanced yield trials.																																				
Activities	:	Continued																																				
Expected outcomes	:	High yield lines with beneficial traits including extra earliness, machine harvesting would be identified.																																				
Observations to be taken	:	Plant height Height of the lowest pod Phenological traits Yield and yield components																																				
Trials	:	Preliminary Yield Trial- for disease resistance and mechanical harvesting- 40 genotypes, two checks with three replication Preliminary Yield Trial –drought tolerant faba bean. 40 genotypes, two checks with three replication Preliminary Yield Trial- heat lines. 40 genotypes, two checks with three replication Preliminary Yield Trial- Large seed types. 40 genotypes, two checks with three replication Preliminary Yield Trial- low tannin trials t (37 genotypes + 2 improved checks + 1 local check) Preliminary Yield Trial – orobanche resistance (40 genotypes + 2 improved checks + 1 local check)																																				
Results		<p>Six preliminary yield trials were conducted at Terbol station Lebanon in Alpha design with three replications. The following traits were recorded: Days to flowering, days to maturity, plant height, and number of nodes until first pods, Grain yield and hundred seed weight. Significant differences among genotypes was presented in (Table 1). Selected lines showed superior yield 10% above the control.</p> <p>Table 1: Variance analysis performed for detecting significance differences of genotypic variation in agronomic, expressed as P-value in all trials.</p> <table border="1"> <thead> <tr> <th>Trial name</th> <th>NG</th> <th>Number of selected lines</th> <th>DFLR</th> <th>DMAT</th> <th>PTHT</th> <th>NNFP</th> <th>Gyha</th> <th>HSW</th> </tr> </thead> <tbody> <tr> <td>Large seeds</td> <td>15</td> <td>5</td> <td>0.001</td> <td>0.001</td> <td>NA</td> <td>0.05</td> <td>0.001</td> <td>0.001</td> </tr> <tr> <td>Low tannin</td> <td>20</td> <td>10</td> <td>0.001</td> <td>0.001</td> <td>0.05</td> <td>0.05</td> <td>0.001</td> <td>0.001</td> </tr> <tr> <td>Machine harvest-diseases</td> <td>40</td> <td>20</td> <td>0.001</td> <td>0.001</td> <td>0.001</td> <td>0.05</td> <td>0.01</td> <td>0.001</td> </tr> </tbody> </table>	Trial name	NG	Number of selected lines	DFLR	DMAT	PTHT	NNFP	Gyha	HSW	Large seeds	15	5	0.001	0.001	NA	0.05	0.001	0.001	Low tannin	20	10	0.001	0.001	0.05	0.05	0.001	0.001	Machine harvest-diseases	40	20	0.001	0.001	0.001	0.05	0.01	0.001
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Machine harvest-diseases	40	20	0.001	0.001	0.001	0.05	0.01	0.001																														

	Heat tolerance	24	12	0.001	0.727	0.397	0.297	0.001	0.001
	Orobanche lines	40	20	0.001	0.001	0.001	0.001	0.001	0.001
	Early and Drought	40	20	0.001	0.001	0.05	NA	0.001	NA
<p>DFLR: days to flowering; DMAT, days to maturity; PTHT, plant height, number of nodes up to first pod (NNFP), GYha, Grain yield per ha, HSW: hundred seed weight.</p>									