

Maize				Sorghum				Rice				Brachypodium			
gene name	location	E-value	score	gene name	location	E-value	score	gene name	location	E-value	score	gene name	location	E-value	score
AC216056.3_FG006	chr. 2: 260449546-260449811	1.30e-96	171	Sb02g037560	71847025-71847612	0	215	Os07g0581700	24220173-24220430	4.00e-81	305	Bradi1g23460.1	18840533-18840785	1.00e-80	303
AC187394.3_FG016	chr. 7: 175520831-175521040	2.50e-54	97	Sb01g043910	67024000-67024247	4.70e-43	139	Os03g0198600	5208766-5209011	2.00e-33	147	Bradi3g35850.1	38113373-38113406	3.00e-07	60
AC195802.3	chr. 1: 31320681-31320864	2.80e-36	81	Sb02g023650	57500046-57500081	1.10e-07	32	Os02g0729700	31269265-31269301	3.00e-11	74	Bradi4g29380.1	34837369-34837401	1.00e-06	58
AC204597.3	107156-107339	2.80e-36	81	chr. 7	52688767-52688802	2.70e-05	28	Os09g0470500	18557594-18557628	1.00e-07	62	Bradi2g06210.1	23246636-23246653	0.073	42
AC197760.3	chr. 1: 26245388-26245444	1.20e-07	33	chr. 4	59114807-59114838	0.006	24	Os08g0481400	23892602-23892636	1.00e-07	62				
AC177946.2	chr. 1: 91646852-91646908	1.20e-07	33	Sb10g029260	59080215-59080234	0.017	20	Os10g0377300	12299374-12299406	2.00e-06	58				
AC194663.3	chr. 2: 222290690-222290724	5.40e-07	31	Sb10g002740	2407492-2407514	0.025	46	Os06g0140400	2123191-2123214	0.002	48				
AC214476.2	chr. 1: 275963972-275964006	1.80e-06	31					chr. 12	21115153-21115175	0.007	46				
AC214506.2	chr. 1: 92405829-92405885	2.80e-05	29					Os04g0660500	34117439-34117459	0.1	42				
AC206905.3	chr. 7: 134611723-134611754	3.10e-05	28												
AC203851.3	chr. 7: 134368691-134368722	0.00011	28												
AC201765.4	chr. 2: 235629943-235629974	0.00011	28												
AC194676.3	chr. 1: 173593361-173593396	0.00011	28												
AC208890.3	chr. 9: 34603456-34603481	0.0017	26												
AC187915.3	chr. 4: 211608369-211608401	0.0069	25												
AC194188.3	chr. 4: 211407014-211407046	0.0069	25												
AC185262.2	chr. 4: 132223151-132223174	0.0019	24												
AC203770.3	chr. 7: 154504042-154504063	0.094	22												

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AC216056.3_FG006	chr. 2: 260449546-260449811	1.30e-96	171	Sb02g037560	71847025-71847612	0	215	Os07g0581700	24220173-24220430	4.00e-81	305	Bradi1g23460.1	18840533-18840785	1.00e-80	303
AC187394.3_FG016	chr. 7: 175520831-175521040	2.50e-54	97	Sb01g043910	67024000-67024247	4.70e-43	139	Os03g0198600	5208766-5209011	2.00e-33	147	Bradi3g35850.1	38113373-38113406	3.00e-07	60
AC195802.3	chr. 1: 31320681-31320864	2.80e-36	81	Sb02g023650	57500046-57500081	1.10e-07	32	Os02g0729700	31269265-31269301	3.00e-11	74	Bradi4g29380.1	34837369-34837401	1.00e-06	58
AC204597.3	107156-107339	2.80e-36	81	chr. 7	52688767-52688802	2.70e-05	28	Os09g0470500	18557594-18557628	1.00e-07	62	Bradi2g06210.1	23246636-23246653	0.073	42
AC197760.3	chr. 1: 26245388-26245444	1.20e-07	33	chr. 4	59114807-59114838	0.006	24	Os08g0481400	23892602-23892636	1.00e-07	62				
AC177946.2	chr. 1: 91646852-91646908	1.20e-07	33	Sb10g029260	59080215-59080234	0.017	20	Os10g0377300	12299374-12299406	2.00e-06	58				
AC194663.3	chr. 2: 222290690-222290724	5.40e-07	31	Sb10g002740	2407492-2407514	0.025	46	Os06g0140400	2123191-2123214	0.002	48				
AC214476.2	chr. 1: 275963972-275964006	1.80e-06	31					chr. 12	21115153-21115175	0.007	46				
AC214506.2	chr. 1: 92405829-92405885	2.80e-05	29					Os04g0660500	34117439-34117459	0.1	42				
AC206905.3	chr. 7: 134611723-134611754	3.10e-05	28												
AC203851.3	chr. 7: 134368691-134368722	0.00011	28												
AC201765.4	chr. 2: 235629943-235629974	0.00011	28												
AC194676.3	chr. 1: 173593361-173593396	0.00011	28												
AC208890.3	chr. 9: 34603456-34603481	0.0017	26												
AC187915.3	chr. 4: 211608369-211608401	0.0069	25												
AC194188.3	chr. 4: 211407014-211407046	0.0069	25												
AC185262.2	chr. 4: 132223151-132223174	0.0019	24												
AC203770.3	chr. 7: 154504042-154504063	0.094	22												