

<b>A</b>	MA	MB	MC	RA	RB	RC
<i>VIT_08s0007g00440</i>	3.77	1.49	23.04	70.10	61.86	1.30
<i>VIT_08s0040g02740</i>	91.72	60.04	79.51	79.33	51.92	58.25
<i>VIT_08s0040g02340</i>	8.83	5.21	3.55	3.47	1.89	1.85
<i>VIT_08s0040g02350</i>	8.92	5.03	3.63	3.34	1.64	2.54
<i>VIT_08s0007g00600</i>	1.28	0.06	0.05	1.18	0.13	0.01
<i>VIT_08s0007g00660</i>	0.57	0.02	0.00	0.86	0.02	0.02
<i>VIT_08s0007g00670</i>	0.42	0.01	0.00	1.41	0.01	0.00
<i>VIT_08s0007g00680</i>	0.70	0.06	0.02	1.46	0.11	0.01
<i>VIT_08s0007g00690</i>	6.82	0.40	0.00	2.54	2.68	0.06
<i>VIT_08s0007g01550</i>	0.04	0.11	0.02	0.26	0.09	0.00
<b>BF</b>	14.13	8.99	6.35	13.56	8.45	9.76

<b>B</b>	MA	MB	MC	RA	RB	RC
<i>VIT_08s0032g01110</i>	68.23	37.26	9.56	90.08	12.00	6.44
<i>VIT_08s0032g00920</i>	93.47	64.25	59.41	50.88	37.58	67.19
<i>VIT_08s0032g01080</i>	6.21	6.25	6.10	3.90	3.53	5.52
<i>VIT_08s0032g01090</i>	8.62	8.51	9.07	5.84	4.59	12.34
<i>VIT_08s0032g01150</i>	44.37	142.86	138.85	32.36	75.99	280.80
<i>VIT_08s0032g01180</i>	25.19	17.05	7.48	18.74	9.92	7.67
<i>VIT_08s0105g00180</i>	5.04	23.72	8.81	1.85	2.44	1.17
<i>VIT_08s0105g00190</i>	3.33	16.72	6.42	2.23	3.63	2.18
<i>VIT_08s0105g00200</i>	0.13	1.55	0.28	0.24	0.23	0.23
<i>VIT_08s0105g00290</i>	27.27	28.17	20.30	25.47	21.68	16.48
<i>VIT_08s0217g00020</i>	90.18	124.60	69.90	100.69	62.54	72.70
<b>ShI</b>	1.37	1.12	1.23	1.19	1.09	1.05

<b>C</b>	MA	MB	MC	RA	RB	RC
<i>VIT_05s0020g01240</i>	13.06	8.02	0.74	6.94	1.19	0.57
<i>VIT_05s0020g02130</i>	94.62	107.50	26.15	133.95	32.12	15.65
<i>VIT_05s0020g01840</i>	6.31	7.88	2.22	17.28	3.19	2.33
<i>VIT_05s0020g03860</i>	3.11	57.11	127.60	1.37	304.66	288.04
<i>VIT_05s0020g03640</i>	1.37	1.99	0.01	6.71	0.35	0.04
<i>VIT_05s0020g03170</i>	10.67	10.97	7.48	58.15	11.73	1.30
<b>MF</b>	0.00	0.00	0.00	0.00	0.00	4.32

Minimum



Maximum